



Significance and occurrence of fumonisins from *Aspergillus niger*

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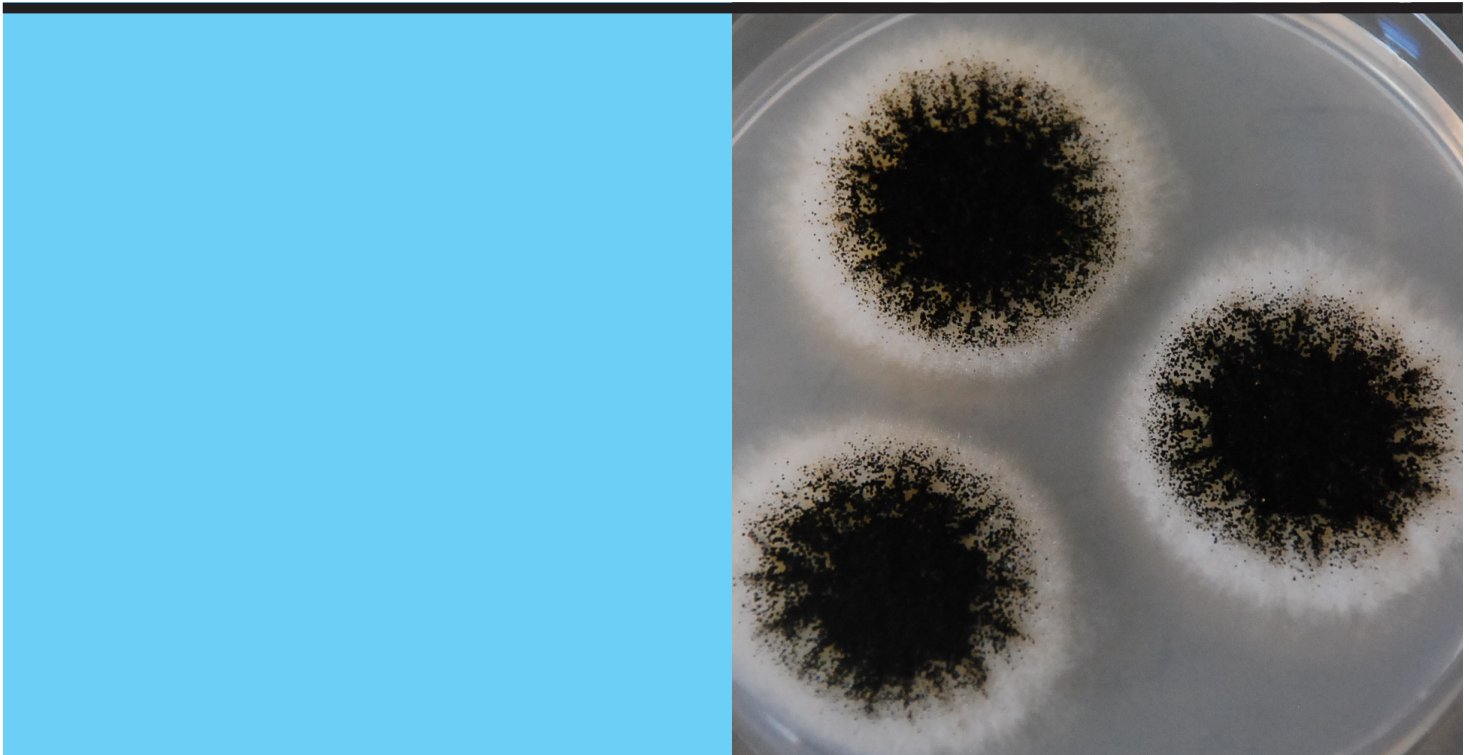
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Significance and occurrence of fumonisins from *Aspergillus niger*



Jesper Mølgaard Mogensen
Ph.D. Thesis
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Preface

This thesis is submitted to the Technical University of Denmark in partial fulfillment of the requirements for the Degree of Doctor of Philosophy in chemistry. The work has been carried out between October 2008 and December 2011, at the Center for Microbial Biotechnology at the Department of Systems Biology under the main supervision of associate professor **Kristian Fog Nielsen** with associate professor **Thomas Ostenfeld Larsen** and Professor **Jens Christian Frisvad** as co-supervisors. The project was funded by the Danish Food Industry Agency (grant 3304-FVEP-07-730-01).

First and foremost I would like to thank my three supervisors for three exciting, interesting and productive years. You have all been very enthusiastic throughout the project and your inspiration scientific advice, constructive criticism and support have been valuable during the PhD study. A thank also goes to **Anoop Kumar Sharma** for his help with the determination of the genotoxicity of fumonisin at DTU Food in Mørkhøj.

I had several opportunities to collaborate with national and international research groups, on various manuscripts. **Rudolf Krska**, **Michael Sulyok** and **Elisabeth Varga** from IFA-Tulln, Austria; **Gordon Shephard** and **Liana van der Westhuizen** from the PROMEC unit, South Africa; **Peter Have Rasmussen** from DTU-FOOD; **Robert Samson** and **Janos Varga** from CBS-KNAW, The Netherlands; **Roman Labuda** from Romer Labs, Austria; **Alena Kubátová** from Charles University, Czech Republic are all thanked for their collaboration and insightful discussions on the various manuscripts included in this thesis.

I would also like to thank **Hanne Jacobsen** and **Jesper Mogensen** for the help with the analytical instrumentation. **Tanja Thorskov Bladt**, **Marie Louise Klejnstrup** and **Lisette Knoth-Nielsen** are acknowledged for their efforts in purifying fumonisins from *A. niger* for the genotoxicity studies. During my PhD project I have had the opportunity to work with several productive students. Most noteworthy are **Stine Mørchholdt Sørensen** for her work on the single kernel manuscript, as well as **Kirsten Amalie Møller** and **Pernille von Freiesleben** for their efforts with *Tolypocladium*.

My current and former office mates, **Maria Månsson**, **Mads Holger Rønne**, **Peter Boldsen Knudsen** and **Tanja Thorskov Bladt** have all been invaluable over the years with your advice, positive attitude and great laughs. I would also like to thank the rest of my colleagues at CMB for contributing to a great work environment where you always feel welcome.

Lastly I would like to thank my family for their support the past three years. I am furthermore grateful for having a great wife and for her invaluable encouragement and support the last 3 years.

Thesis outline

The overall aim of this thesis was to determine whether fumonisins produced by *Aspergillus niger* is an overlooked health risk in Denmark and worldwide. The objectives were to survey the occurrence and levels of fumonisin in the relevant food commodities such as grapes (**paper 2**), wine (**Paper 3**), raisins (**Paper 2+4**), black tea (**Paper 5**), and peanuts (**Paper 6**); investigation of other fungi with similar genes for potential fumonisin production (**Paper 8**); and to determine the genotoxic potential of fumonisins from *A. niger*. Other objectives were to investigate the effect of physiological parameters on the production of fumonisins by *A. niger* (**Paper 1**) and the fumonisin content in single maize kernels (**Paper 7**). The secondary metabolism of black Aspergilli was reviewed in (**Paper 9**).

The first chapter introduces mycotoxins with special focus on black Aspergilli and the toxins they produce, specifically fumonisins. Chapter 2 outlines the experimental setup for the thesis. In chapter 3 the main results of the articles are presented and discussed. Finally, chapter 4 is the overall conclusion.

Summary

Fumonisins is a well-studied group of mycotoxins, mainly produced in maize by *Fusarium* species. However with the recent discovery of a fumonisin production by *Aspergillus niger*, other food commodities are at risk, since *A. niger* is a ubiquitous contaminant of many food and feed products.

The objective of this thesis was to determine the significance and occurrence of fumonisins from *Aspergillus niger* in food, the frequency of fumonisin production in *A. niger* isolates, as well as studies of the effect of physiological factors affecting fumonisin production. Major findings in this context have been the documentation of the production of fumonisins in raisins and peanuts, and occurrence of *A. niger* derived fumonisins in retail wine and raisins.

Physiological investigations have demonstrated that fumonisin production in *A. niger* occurs at temperatures between 20-37 °C. Three water activity lowering solutes, glycerol, NaCl, and sucrose were used. All *A. niger* strains responded differently on the three solutes. Addition of NaCl and sucrose increased the fumonisin production, where addition of glycerol decreased the production. Fumonisin production in grape and derived products was investigated. Sixty-six *A. niger* strains were isolated and screened for a fumonisin production, and 77 % produced fumonisin on agar media. Worst-case scenarios were made by inoculation of *A. niger* on grapes and solidified the thought that a potential fumonisin production/contamination in grapes and derived products could occur. All tested *A. niger* strains produced fumonisins, not only when growing on grapes, but also on dried grapes and re-moistened raisins. By screening of 77 wine and 21 raisins samples, 18 wine (1-25 µg FB₂/l) and 10 raisin (1.3-13 µg FB₂/kg) samples contained detectable amounts of fumonisins.

An investigation of peanuts and peanut butter did not show any sign of fumonisin contamination, even though *A. niger* produced high amounts of fumonisins in worst case scenarios on shelled peanuts. Black and puerh tea were also examined, since *A. niger* were claimed to be the primary black Aspergilli in these. After isolation of 47 black Aspergilli, who all turned out to be *A. acidus* and did not produce neither ochratoxin or fumonisins, no significant risk from these toxins were associated with this intake.

Single kernel analysis of maize showed that fumonisins were present in 15% of the kernels at varying concentration. The frequency of highly contaminated kernels was 4 %. An estimated

calculation showed that the fumonisin content could be reduced with more than 70 % by sorting the maize. Other frequently found fungal metabolites in the maize kernels were chanoclavine, emodin, equisetin, and fusaric acid.

The biotechnologically important fungus *Tolypocladium*, a producer of the immunosuppressive drug cyclosporin, was identified as a new fumonisin producer. Three species within the genera produced fumonisin B₂ and B₄, and due to the widespread ability within the genus, it should be determined if this mycotoxin production occurs under industrial settings.

Dansk sammendrag

Fumonisin er en gruppe velstuderede mykotoksiner, der hovedsagligt produceres af *Fusarium* i majs. Med opdagelsen af, at *Aspergillus niger* kan producere fumonisiner, er der flere grupper fødevarer som kan indeholde fumonisin, da *A. niger* er en hyppig kontaminant på mange forskellige fødevarer.

Formålet med denne Ph.D. afhandling var at undersøge udbredelsen og betydningen af *A. niger*'s produktion af fumonisiner i fødevarer, frekvensen af fumonisin producerende *A. niger*, samt effekten af fysiologiske faktorer på fumonisin produktionen. De væsentligste opdagelser, er fund af fumonisiner i indkøbte vine og rosiner, samt dokumentation af høj fumonisin produktion i kunstigt inficerede rosiner og jordnødder.

Fysiologiske undersøgelser viste, at *A. niger* producerer fumonisin ved temperaturer mellem 20-37 °C. Til at sænke vandaktiviteten blev NaCl, Glycerol og sucrose anvendt. Alle *A. niger* stammer reagerede forskelligt på de tre forskellige typer af medier med sænket vandaktivitet. Tilsætning af lave mængder NaCl og sucrose forøgede fumonisin produktionen, mens tilsætning af glycerol formindskede den.

Produktionen af fumonisiner i fødevarer herunder druer og afledte produkter blev også undersøgt. Seksogtres *A. niger* stammer blev isoleret fra rosiner og screenet for fumonisin produktion. I alt producerede 77 % af stammerne fumonisin på agar medier. *A. niger* blev podet på druer for at efterligne værste tænkelige scenarier. Da samtlige *A. niger* producerede fumonisin under disse forhold, var der mulighed for at fumonisin kunne findes i drue afledte produkter. Alle de testede stammer producerede desuden fumonisin i tørrede druer og fugtede rosiner. Syvoghalvfjerds vine og 21 rosin mærker blev undersøgt for fumonisiner og 18 vine (1-25 µg FB₂/l) samt 10 rosin mærker (1.3-13 µg FB₂/kg) indeholdt målbare mængder.

Undersøgelsen af fumonisiner i jordnødder og jordnøddesmør, påviste ikke tilstedeværelsen af disse mykotoksiner på trods af, at *A. niger* kan producere fumonisin i værste tænkelige scenarier i jordnødder. Sort og Puerh te blev også undersøgt, da det var påstået, at *A. niger* var den dominerende organisme fra section *Nigri*. Men efter at have isoleret 47 sorte aspergiller, som alle blev identificeret som *A. acidus*, der ikke producerede hverken ochratoxin A eller fumonisiner, var der ikke tegn på en signifikant risiko ved indtagelse af te.

VII | Dansk sammendrag

Enkelt kerne analyse af majs påviste, at fumonisin fandtes i 15 % af kernerne med varierende koncentration. Frekvensen af kerner med en høj koncentration af fumonisin var 4 %. Hvis disse højt inficerede kerner kan frasorteres vil fumonisinindholdet i majsene reduceres med 70 %. Øvrige hyppige svampe metabolitter detekteret i majs kernerne var chanoclavine, emodin, equisetin, samt fusaric acid.

Den bioteknologiske anvendte skimmelsvamp *Tolypocladium*, der anvendes til produktion af det immunosuppressive medikament cyclosporin, blev identificeret som en ny fumonisin producent. Tre arter i denne slægt producerede fumonisin B₂ og B₄ og på grund af den industrielle brug af denne slægt, bør det undersøges om disse mykotoksiner produceres under industrielle betingelser.

List of original papers

- Paper 1:** J.M. Mogensen, K.F. Nielsen, R.A. Samson, J.C. Frisvad and U. Thrane. Effect of temperature and water activity on the production of fumonisins by *Aspergillus niger* and different *Fusarium* spp. BMC Microbiology. 2009. 9:281
- Paper 2:** J.M. Mogensen, J.C. Frisvad, U. Thrane, K.F. Nielsen. Production of fumonisin B₂ and B₄ by *Aspergillus niger* on grapes and raisins. Journal of Agricultural and Food Chemistry. 2010, 58:954-958
- Paper 3:** J.M. Mogensen, T.O. Larsen, K.F. Nielsen. Widespread occurrence of the mycotoxin Fumonisin B₂ in wine. Journal of Agricultural and Food Chemistry. 2010, 58:4583-4587
- Paper 4:** P. B. Knudsen, J. M. Mogensen, T. O. Larsen and K. F. Nielsen. Occurrence of fumonisins B₂ and B₄ in retail raisins. Journal of Agricultural and Food Chemistry. 2011, 59:772-776
- Paper 5:** J.M. Mogensen, J. Varga, U. Thrane and J.C. Frisvad. *Aspergillus acidus* from Puerh tea and black tea does not produce ochratoxin A and fumonisin B₂. International Journal of Food Microbiology. 2009. 132:141-144.
- Paper 6:** J. M. Mogensen, E. O. Søndergaard and K. F. Nielsen. Studies on fumonisins in peanuts and peanut butter. Journal of Agricultural and Food Chemistry. Submitted.
- Paper 7:** J. M. Mogensen, S. M. Sørensen, M. Sulyok, L. van der Westhuizen, G. Shephard, J.C. Frisvad, U. Thrane, R. Krska, K. F. Nielsen. Single kernel analysis of fumonisins and other fungal metabolites in maize from South African subsistence farmers. Food Additives & Contaminants. 2011. 28:1724-1734.
- Paper 8:** J. M. Mogensen, K. A. Møller, P. von Freiesleben, R. Labuda, E. Varga, M. Sulyok, A. Kubátová, U. Thrane, B. Andersen, K. F. Nielsen. Production of Fumonisin B₂ and B₄ in *Tolypocladium* species. Journal of Industrial Microbiology and Biotechnology. 2011, 38:1329-1335.
- Paper 9:** K.F. Nielsen, J.M. Mogensen, M. Johansen, T.O. Larsen and J.C. Frisvad. Review of secondary metabolites and mycotoxins from the *Aspergillus niger* group. Analytical & Bioanalytical Chemistry. 2009. 395:1225-1242.

IX | List of original papers

Other papers (non peer review) in Danish and conference proceedings

J.M. Mogensen, J. Mogensen, J.C. Frisvad, U. Thrane, T.O. Larsen and K.F. Nielsen.
Giftige fumonisiner fra *Aspergillus niger*. Dansk Kemi. 2009, 90(1):25-27

J. M. Mogensen, P. B. Knudsen, T. O. Larsen, J. C. Frisvad, U. Thrane and K. F. Nielsen. Fumonisin from *Aspergillus niger* in grapes and derived products. Poster at Mycored Africa 2011 (4th-6th April), Cape Town, South Africa

J. M. Mogensen, S. M. Sørensen, M. Sulyok, G. Shephard, L. van der Westhuizen, J. C. Frisvad, U. Thrane, K. F. Nielsen. Fumonisin and other compounds in South African maize—A single kernel approach. Poster at Mycored Africa 2011 (4th-6th April), Cape Town, South Africa

J. M. Mogensen, Peter B. Knudsen, Jens C. Frisvad, Thomas O. Larsen and Kristian F. Nielsen. Fumonisin from *Aspergillus niger* in grapes and derived products. Poster at 32nd Mycotoxin Workshop 2010 (14th-16th June), Lyngby, Denmark.

J.M. Mogensen, J.C. Frisvad, T.O. Larsen and K.F. Nielsen. Fumonisin and other compounds from *Aspergillus niger*. How can an organism be both an industrial workhorse and a food safety problem? Presentation at the 31st Mycotoxin Workshop 2009 (15th-17th June), Münster, Germany

J. M. Mogensen, U. Thrane, J. Varga, T. O. Larsen, J. C. Frisvad and K. F. Nielsen. Fumonisin from *Aspergillus niger* – A new health risk? Poster at Fungi and Health Symposium, Amsterdam, the Netherlands 2008

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1 Introduction

1.1 Fungi & Mycotoxins

Fungal contamination of crops has been a known problem since ancient times. In the 700-800 BC Romans made sacrifices to the god Robigus who had to be appeased in order to protect the grain and trees^[1]. The presence of fungi in the field is also mentioned several times in biblical literature, e.g.: “Many times I struck your gardens and vineyards, destroying them with **blight** and **mildew**. Locusts devoured your fig and olive trees, yet you have not returned to me, declares the LORD” (Amos 4:9).

Mycotoxins are secondary metabolites produced by filamentous fungi. Mycotoxins are a diverse range of natural products originating from different biosynthetic pathways. The most common definition of mycotoxins is: “Mycotoxins are natural products produced by fungi that evoke a toxic response when introduced in low concentration to higher vertebrates and other animals by a natural route.”^[2,3]. Mycotoxins are distributed worldwide in food and feed products, and are formed under certain environmental conditions e.g. proper moisture, suitable temperature, physical damage to the commodity, and presence of the fungal spores^[4-7]. Mycotoxins are produced pre-harvest in the field as well as during postharvest transportation and/or storage^[4,5,7,8], causing severe economic losses due to e.g. poor crop quality, yield loss, impaired animal health, and higher mortality rates^[9]. As a consequence export of several agricultural commodities, especially crops originating from developmental countries is affected. Good agricultural practices are therefore essential to avoid mycotoxin contamination, because poor harvesting practices, improper drying, handling, packaging, storage, and transport conditions increase fungal growth and mycotoxin production^[5].

Diseases caused by mycotoxins, mycotoxicoses, have been known since ancient times and several interpretations have linked historical events to the ingestion of mycotoxins^[10-13]. For instance the involvement of mycotoxins in the plague of Athens in 430 BC^[14], episodes occurring in the Middle Ages, referred to as “St. Anthony’s Fire”^[13] and the mycotoxin T-2 toxin as a suspected causative agent for numerous alimentary toxic aleukia related fatalities in Russia during World War II^[15]. The fundamental moment for modern mycotoxicology was however, the outbreak of Turkey-X disease in the United Kingdom in 1960, which caused the death of thousands of turkeys and ducklings, and was later attributed to be caused by aflatoxins^[16-19]. This event is seen as the cause for the massive

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expansion of research efforts in the areas of mycotoxicology, such as mode of action, implication for human health, and chemical determination of new toxins^[8]. Based on this huge effort within fungal pathogens of major crops, their natural occurrence and implications for human health, researchers has focused mainly on aflatoxins, fumonisins, trichothecenes, zearalenone, ergot alkaloids, ochratoxin A, and patulin^[5,8], figure 1. Although hundreds of fungal toxins are known, a more limited number are generally considered to play an important role in food safety.

Mycotoxins of concern are mainly produced by the three genera *Aspergillus*, *Fusarium* and *Penicillium*^[20].

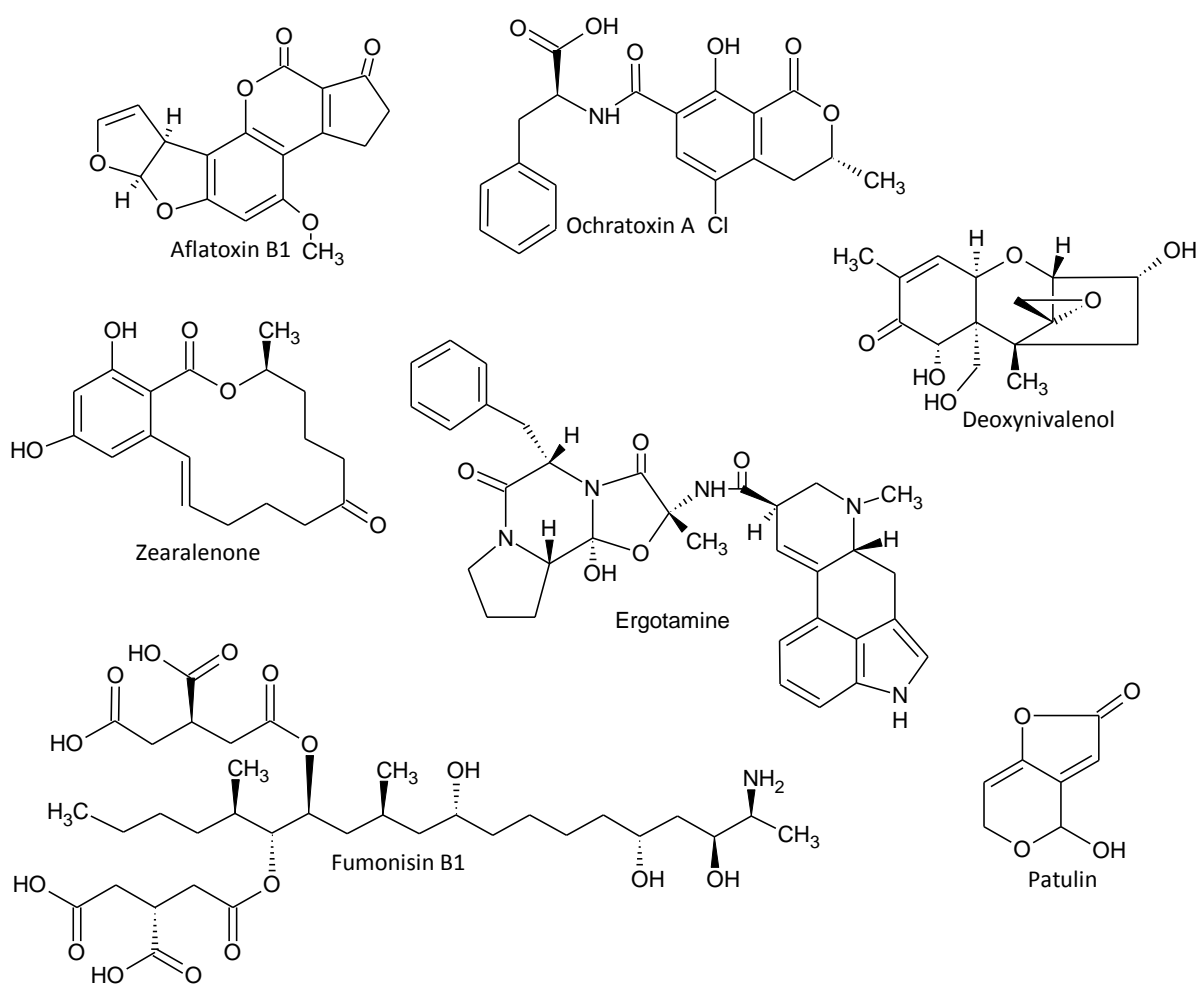


Figure 1: Common regulated toxins.

Within the group of mycotoxins, aflatoxins, ochratoxins, and fumonisins are considered the most important. All three types of mycotoxins are highly regulated, due to their toxicity and prevalence^[21]. Aflatoxins are acutely toxic, immunosuppressive, mutagenic, teratogenic and

carcinogenic compounds^[4,22]. Aflatoxins are produced by *Aspergilli*^[16,19], and are often associated with maize and peanuts^[23,24]. According to USFDA the advisory limit for aflatoxins is 50 µg/kg in all food types (except milk) and EU has regulatory limits of 0.05-15 µg/kg depending on the food type^[21,25]. Ochratoxin A has been shown to be nephrotoxic, immunosuppressive, carcinogenic and teratogenic^[4,22]. Ochratoxin A are produced by *Penicillia* and *Aspergilli*^[26,27] and frequently found in cereals, grapes (derived products) and coffee beans^[5,23]. The regulatory limits for ochratoxin A are 2-10 µg/kg according to EU^[21,28]; however USFDA has not set any advisory limits for ochratoxin A^[29,30]. Fumonisins are cardiotoxic, hepatotoxic, nephrotoxic and associated with several human and animal diseases^[31] (elaborated in chapter 1.2.1). Fumonisins are mainly produced by *Fusaria* (primarily section *Liseola*)^[32,33], but *Aspergillus niger*^[34-36] is also reported as a producer. Fumonisins are primarily a concern in maize and the EU has set maximum levels of 0.2-2 mg/kg^[37] for fumonisins where the USFDA has recommended a maximum levels of 2-4 mg/kg^[38]. Regulated and examples of non regulated mycotoxins, their associated commodities along with regulatory levels are given in table 1

Table 1: Examples of mycotoxins; their source, commodities and regulatory limit in food^[5,8,13,20,21,23,39,40]

Mycotoxin	Primary source	Important commodities	Regulatory limits in food (µg/kg)	
			USFDA	EU
Aflatoxin	<i>A. flavus</i> , <i>A. parasiticus</i>	Maize, peanuts	20	0.01-15
Alternariol	<i>Alternaria tenuissima</i>	Cereals	-	-
Citrinin	<i>P. citrinum</i> , <i>P. expansum</i>	Grain	-	-
Ergot alkaloids	<i>Claviceps purpurea</i>	Grain, primarily rye	-	1000000 (feed)
Fumonisins	<i>F. verticillioides</i> , <i>F. proliferatum</i>	Maize	2000-4000	200-4000
Gliotoxin	<i>A. fumigatus</i>	Virulence factor in mammalian mycoses	-	-
Ochratoxin	<i>A. carbonarius</i> , <i>A. ochraceus</i> , <i>P. verrucosum</i>	Cereals, grapes, coffee beans	-	Food: 0.5-80
Patulin	<i>P. expansum</i>	Apples, apple juice	50	10-50
Sterigmatocystin	<i>A. nidulans</i> , <i>A. versicolor</i>	Moldy grain, green coffee beans and cheese	-	-
Deoxynivalenol	<i>F. equiseti</i> , <i>F. graminearum</i>	Cereals	1000	200-1750
Zearalenone	<i>F. graminearum</i> , <i>F. culmorum</i>	Cereals	-	20-400

1.2 Fumonisins

The fumonisins were first isolated from *Fusarium verticillioides* (= *Fusarium moniliforme*) by Bezuidenhout *et al.* in 1988^[33]. Analogues from *Fusarium* have been identified since and are classified in four structural series of fumonisin designated A, B, C, P (figure 2)^[33,41,42]. The four

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groups have the same scaffold but differ in selected structural moieties. The A-groups differ from the B-group by the amine which is acetylated, the C-group consist of a C19 backbone instead of C20, the P-series have a 3-hydroxypyridinium group instead of the amino group at position C2 [33,41,42]. The B-series is the most abundant naturally occurring fumonisins with FB₁ as the predominant^[32,43].

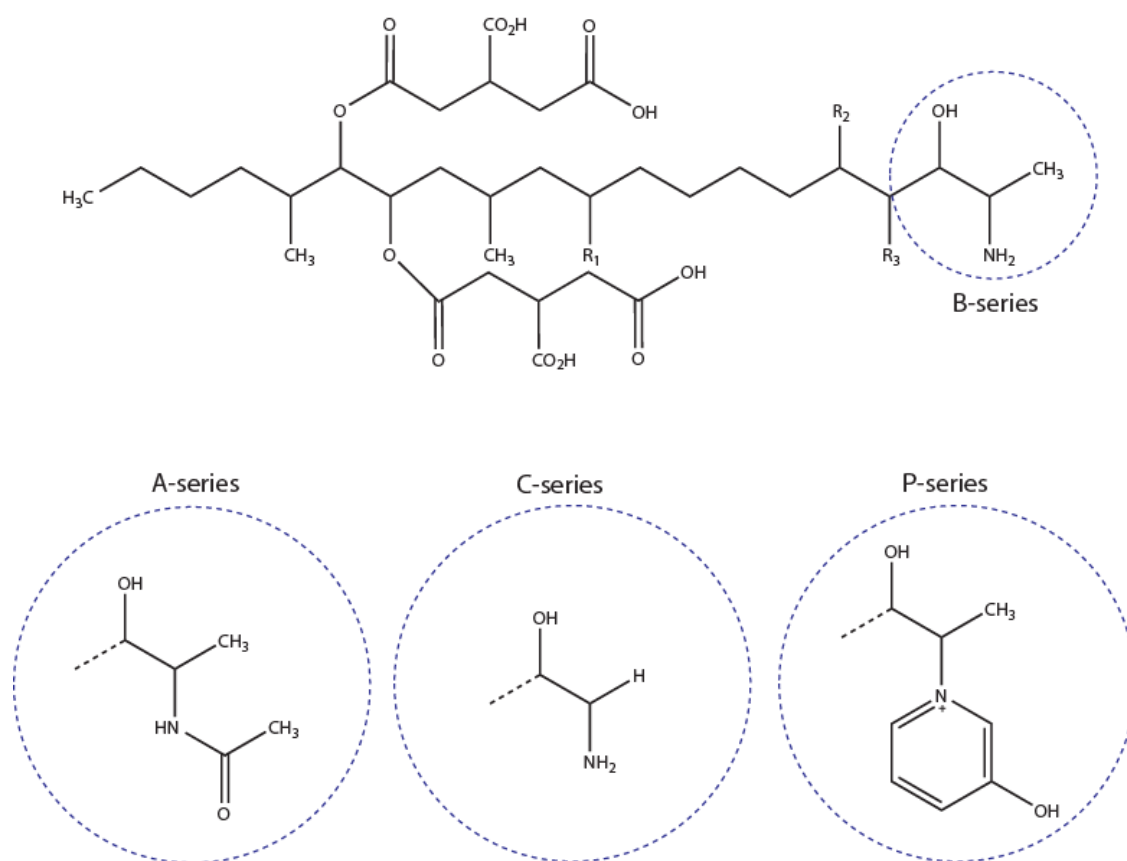


Figure 2: Overall structure of fumonisins A, B, C, P-series.

The event that led to the discovery of the fumonisins was a field outbreak of equine leukoencephalomalacia (ELEM) in South Africa in 1970^[44]. During the investigation of this event, *F. verticillioides* was isolated from a batch of moldy maize believed to be the causative agent of the ELEM outbreak^[45]. Focusing the research on this target organism (*F. verticillioides*), showed that a number of these caused ELEM in horses, pulmonary edema in pigs and was highly hepatotoxic and cardiotoxic in rats^[46,47]. In 1988, a breakthrough came with the isolation of FA₁, FA₂, FB₁, and FB₂^[33]. In the following years, several *Fusarium* species were identified as fumonisins producers^[32,43,48] as

well as identification and structural elucidation of numerous fumonisins analogs^[42,49,50]. The fumonisin producing *Fusaria* are primarily found within section *Liseola*, mainly *F. proliferatum* and *F. verticillioides*^[32], however species of section *Discolor*, *Elegans*, *Gibbosum*, and *Martiella* are also reported as fumonisin producers^[34,48,51]. A single report has furthermore claimed that *Alternaria arborescens* (reported as *A. alternata*) produced FB₂, but this strain has since lost this ability^[52-54]. In 2006, a surprising discovery was made, when full genome sequencing of *A. niger* showed that the genome included a putative *Fum* gene cluster^[55,56]. This gene cluster could be silenced and hereby not expressed, but in 2007 it was shown that *A. niger* is able to produce FB₂, and this was followed by the discovery of production of FB₄ and the new FB₆^[34-36]. In total fumonisin production has been reported in 26 *Fusarium*, one *Alternaria* and one *Aspergillus* species, table 2.

Table 2: Species reported as fumonisin producers and their production of fumonisins

Species	Fumonisins	References
<i>Fusarium acutatum</i>	B ₁ , B ₂	[57]
<i>F. andiyazi</i>	B ₁ , B ₂	[58-60]
<i>F. anthophilum</i>	B ₁ , B ₂	[51,61]
<i>F. begoniae</i>	B ₁	[57]
<i>F. brevicatenilatum</i>	B ₁	[57]
<i>F. dlamini</i>	B ₁	[51]
<i>F. equiseti</i>	B ₁	[48]
<i>F. fujikuroi</i>	B ₁ , B ₂ , B ₃	[60,62,63]
<i>F. globosum</i>	B ₁ , B ₂ , B ₃	[62,64]
<i>F. lactis</i>	B ₁	[65]
<i>F. napiforme</i>	B ₁	[51]
<i>F. nygamai</i>	A ₁ , A ₂ , A ₃ , B ₁ , B ₂ , B ₃ , B ₄ , B ₅ , C ₁	[43,51,58,66,67]
<i>F. oxysporum</i>	B ₁ , B ₂ , B ₃ , C ₁ , C ₄	[41,48,68]
<i>F. phyllophilum</i>	B ₁	[57]
<i>F. polyphialidicum</i>	B ₁	[69]
<i>F. proliferatum</i>	A ₁ , A ₂ , A ₃ , B ₁ , B ₂ , B ₃ , C ₁ , C ₄ , P ₁ , P ₂ , P ₃	[51,66-68]
<i>F. pseudocircinatum</i>	B ₁ , B ₂	[57]
<i>F. pseudonygamai</i>	B ₁ , B ₂	[58]
<i>F. ramigenum</i>	B ₁ , B ₂	[70]
<i>F. redolens</i>	B ₁ , B ₂ , B ₃	[71]
<i>F. sacchari</i>	B ₁	[59,72]
<i>F. sambucinum</i>	B ₁	[73]
<i>F. solani</i>	B ₁	[73]
<i>F. subglutinans</i>	B ₁ , B ₂ , B ₃	[63,72]
<i>F. thapsinum</i>	B ₁ , B ₂ , B ₃	[58,59,74]
<i>F. verticillioides</i>	A ₁ , A ₂ , A ₃ , B ₁ , B ₂ , B ₃ , B ₄ , B ₅ , C ₁ , C ₃ , C ₄ , P ₁ , P ₂ , P ₃	[33,58,66,68]
<i>Alternaria arborescens</i> *	B ₁	[52]
<i>Aspergillus niger</i> (and <i>A. awamori</i>)	B ₂ , B ₄ , B ₆	[34,35]

*reported as *Alternaria alternata* (Fr.) Kiessl. f. sp. *lycopersici*. This strain has since lost its ability to produce FB₁

1.2.1 Toxicity of fumonisins

Numerous toxicity studies of fumonisins have been performed, using various assays to determine the different types of toxicity. This is important due to the link to both human and animal toxicoses from consumption of contaminated maize-based food and feeds^[75]. As the most abundant fumonisin, FB₁ is the most thoroughly investigated concerning the toxicity of fumonisins. The causality between fumonisins and animals has been proved in several cases^[31,76]. Fumonisin and *F. verticillioides* contaminated feeds have been recognized as the cause for the neurotoxic disease ELEM in horses^[77,78], and are furthermore cardiotoxic and cause pulmonary edema in pigs^[79]. Cattle and poultry appear to be less sensitive than horses, pigs, rabbits and laboratory rodents^[31], resulting in a less strict guidance levels in feed specified for these^[38]. It has been shown in laboratory animals that liver and kidney are the major target organs; however, species-, strain- and sex-dependent differences in dose response occur^[31,76,80]. The effect of fumonisins on human health is uncertain^[76]. Epidemiologic studies around the world have revealed that the incidence of human esophageal cancer and the occurrence of *F. verticillioides* appear to correlate^[31]. Furthermore, fumonisins are suspected risk factors for liver cancer^[81,82], neural tube defects^[83-86], and cardiovascular problems^[87] in populations with a high intake of food made with contaminated maize.

Concerning the genotoxic potential of fumonisins, several independent studies have been performed; however, with conflicting results, since reports of positive and negative results in various assays (both *in vivo* and *in vitro*) have been published^[31].

Fumonisin are structurally similar to the sphingolipids precursors^[88]. These similarities cause a disruption/inhibition of the sphingolipid biosynthesis, specifically the ceramide synthase^[88]. It is particular fumonisins with an amino group at C2, which competitively inhibit ceramide synthase, and hereby disrupt the *de novo* biosynthesis of ceramide and sphingolipid metabolism^[89,90].

Inhibition of biosynthesis of sphingolipids is seen at different levels, and is reflected in changes of the ratio sphinganine/sphingosine (Sa/So)^[91]. Although the specific effects of sphingolipids on apoptosis are complex and not fully understood, there is no doubt that in various *in vivo* and *in vitro* cellular systems, administration of FB₁ produces alterations in sphingolipids with consequent alterations of apoptosis and cell proliferation^[92]. The results suggesting that the amino group appears to be the reason for the ceramide synthase inhibition and toxicity of fumonisins, appear

to be true since N- substituted fumonisins showed reduced toxicity both *in vivo* and *in vitro*^[93-95] and less ceramide synthase inhibiting ability^[96].

Since the molecular structures of FB₂ and FB₃ are almost identical to FB₁, it has been claimed that the degree of toxicity may be at the same level as that of FB₁^[94,97,98]. From a *Fusarium* point of view, the importance of their toxicity is marginal, since the levels of these compounds are usually 4-10 times lower than FB₁. However with the production of FB₂, FB₄ and FB₆ by *A. niger* further research about these compounds is necessary.

1.2.2 Analysis of fumonisins in food & feed

The widespread prevalence and toxicity of fumonisins resulted in the development of a number of analytical methods for detection of fumonisin levels in food and feeds, table 3. In the first report of the isolation of fumonisins, Gelderblom *et al.* used normal phase chromatography as well as HPLC^[99]. Since then, the need for quantitative analytical methods for detection of fumonisins has been obvious. Initially the primary focus was on gas chromatography^[100-103]. GC-MS analysis is based on hydrolysis of the esterified side chain and trimethylsilyl or trifluoroacetate derivatization, and is sensitive and selective, but at the time involved expensive equipment^[104].

Table 3: Detection methods used in fumonisin analysis and examples of their advantages/disadvantages

Detection method	Chemical modification	Advantages	Disadvantages
HPLC-FLD	Derivatization of the amine	sensitive, widespread , equal response factor, low initial cost	Derivatives decompose quickly leading to underestimation, derevatization of every amine group, high running costs, clean-up procedures
GC-MS (rare)	Hydrolysis and esterification	Sensitive, selective	Complicated pre-treatment, multiple handling steps
TLC	Ninhydrin, <i>p</i> -anisaldehyde or flourescamine spray	Cheap, simple, rapid	Lack of automation, sensitivity, Verification by GC or LC
ELISA	None	Easy, rapid, inexpensive, Sensitive, specific	Overestimates fumonisin level, less sensitive than LC, cross reactivity, Verification by GC or LC
LC-MS	None	No chemical modification, less sample prep than other, crude extract can be analyzed, high sensitivity	Expensive equipment, ion-suppression

However focus shifted towards HPLC methods, due to the multiple handling steps (hydrolysis, clean-up, derivatization) prior to GC-MS analysis^[105]. Since HPLC methods were easier to apply and required less sophisticated equipment, they became the method of choice for detection and quantification, primarily HPLC coupled to a fluorescence detector (FLD)^[105,106]. The fumonisins are highly polar molecules (logD -1.5 at pH 3.2 and -4.21 at pH 7) which are soluble in water and polar solvents, and are suited for separation by reversed phase HPLC^[105] (acidic). Since fumonisins lack UV absorption, and therefore undetectable with UV or FLD, chemical modification like derivatization of the amine with o-phthaldialdehyde (OPA) is necessary to make it detectable^[107]. However the disadvantages of OPA derivatives of fumonisins are, that they decompose very rapidly, thus leading to underestimation of the actual level of contamination^[108]. Additionally, OPA may react with other amino group containing compounds, resulting in some interference^[108]. Chromatographic detection has mostly been achieved with a UV and fluorescence detector, advances within liquid chromatography–mass spectrometry (LC–MS) technology have however, shifted the emphasis towards this analytical approach^[105,108,109]. The LC-MS provides quantitative analyses as well as confirmation of the identity of the fumonisins without any need of derivatization. Moreover, the commercial availability of the isotopically labeled FB₁ allowed the development of LC/MS methods based on the isotopic dilution approach, which are usually characterized by high accuracy^[108,110]. Furthermore virtually all mycotoxin analytes are compatible with LC-MS because of the conditions applied during separation and detection^[109,110]. Electro Spray Ionisation (ESI)-MS is an ideal way to ionize, detect, and measure fumonisins, which are ionic compounds and therefore produce abundant signals in both positive and negative ion modes, however positive ion mode is most commonly used^[108,110,111]. Compared to the other detection techniques, the major advantage of the LC/ESI/MS detection of fumonisins is beside that no derivatization or special sample preparation is required, no fragmentation is produced during the ionization^[108,110,111]. For these reasons, it is possible to directly analyze culture materials and food products by simply extracting, filtering, and injecting the sample onto the LC column. Additionally, a recent comparison of fumonisin detection using HPLC-FLD, LC-MS and LC-MS/MS was reported, proving that LC-MS/MS had the highest sensitivity followed by HPLC-FLD and LC-MS^[112].

Immuno based techniques such as ELISA have also been used for detection of fumonisins in food and feed^[8,105,107,109,110]. The advantages of this approach is that it is easy, rapid and

inexpensive^[107], however ELISA methods generally overestimate the fumonisin content in samples and positive samples needs verification by other means^[107,108].

Often the crude extracts must be purified in order to remove matrix components and to concentrate the fumonisins prior to analytical chromatographic analysis. Examples of sample clean-up with different approaches, involved the usage of reversed phase, strong anion-exchange (SAX) or immunoaffinity columns (IAC)^[113,114]. IAC provides the most selective purification, and the user obtain cleaner extracts with a minimum of interfering matrix components and leading to excellent signal-to-noise ratios compared to the use of the less selective SPE sorbent materials^[110]. However, IAC have a lower binding capacity (1-2 µg FB's) and therefore IAC is not recommendable for purification and determination of high amounts of FB^[105,107]. SAX columns have showed a superior purification compared to reversed C18 phase columns^[115]; however, SAX require a pH of the sample extract above 5.8 for adequate retention^[116].

1.2.3 Fumonisins in Food & Feed - Recent findings

Fumonisins are found frequently in maize, and almost all reported findings during the last 30 years are reported in maize^[3,31,117,118]. The reported concentrations vary a great deal^[3,117], with levels up to 300 mg/kg observed in Italy^[31,119]. Although fumonisins in maize and derived products are the primary problem, reports of fumonisins in other commodities have been published, for instance rice^[120], sorghum^[121], black tea^[122] and beer^[123]. Table 4 illustrates examples of recent findings of fumonisins in food commodities other than maize and maize products.

Table 4: Examples of recent findings (2005-2011) of fumonisin B₁ associated with *Fusarium* in non-maize products

Food type	Country	Frequency	Levels (µg/kg)	References
Asparagus	China	80% (n=80)	24-670	[124]
Beans	Cameroon	20% (n=15)	28-1351	[125]
Beer	European	97% (n=33)	0.1-30.3	[126]
Cereals	Malaysia	19% (n=100)	41.3-209.3	[127]
Dried figs	Turkey	75% (n=15)	0-3649	[128]
Milk	Italy	80%	0.23-0.46	[129]
Oats	Argentina	9% (n=23)	105-108	[130]
Peanuts	Cameroon	19% (n=16)	25-1498	[125]
Retail food	Japan	23% (n=810)	0-1380	[131]
Rice	Canada	15%	4.5 (Mean)	[132]
Sorghum opaque beers	Cameroon	88% (n=120)	0.0-230	[133]
Spices	Tunisia	15% (n=13)	70-130	[134]
Tea	China/Belgium	1% (n=91)	76	[135]

With the recent finding of a fumonisin production in *A. niger*^[34-36], new food commodities are at risk. Until now there are very few reports of fumonisins associated with *A. niger* in food and feed, these are shown in table 5. A general trend of these fumonisin levels are that levels associated with *A. niger* are markedly lower than levels in commodities associated with *Fusarium*.

Table 5 Findings of fumonisin B₂ associated with *Aspergillus niger* in food.

Food type	Country	Frequency	Levels (µg/kg)	References
Raisin	6 different countries	54% (n=13)	157-11516	[136]
Grape must	Italy	15% (n=12)	10-400	[137]
Wine	Italy	18% (n=51)	0.4-2.4	[114]
Coffee beans	Thailand	58% (n=12)	1.3-9.7	[36]

1.3 Black Aspergilli

1.3.1 Occurrence & taxonomy

The genus *Aspergillus* consists of more than 250 species^[138]. Within the genus are the black Aspergilli (*Aspergillus* subgenus *Circumdati* section *Nigri*), which are an important group in food mycology and biotechnology. Members of section *Nigri* are ubiquitous and have been isolated from a wide variety of substrates found around the world^[139-143]. The section *Nigri* consists of 25 species including both uniserate and biserate species (table 6) of which *A. acidus*, *A. brasiliensis*, *A. carbonarius*, *A. ibericus*, *A. niger*, and *A. tubingensis* are found commonly^[144-146]. The rest of section *Nigri* are rare and each species is suggested to represent a frequency below 1%^[146]. Within section *Nigri*, *A. niger* is the most common contaminants of food^[146], especially fruits and certain vegetables^[143], and is frequently determined in grapes, green coffee beans, onions, mango, maize and other cereals, peanuts, dried fruits, and many other products^[36,143,147].

Table 6: Overview of *Aspergillus* section *Nigri*. (Bold: Most common)^[144,145]

Clade	Species
<i>A. niger</i> clade	<i>A. acidus</i> , <i>A. awamori</i> , <i>A. brasiliensis</i> , <i>A. costaricensis</i> , <i>A. eucalypticola</i> , <i>A. neoniger</i> , <i>A. niger</i> , <i>A. piperis</i> , <i>A. tubingensis</i> , <i>A. vadensis</i>
<i>A. carbonarius</i> clade	<i>A. carbonarius</i> , <i>A. ibericus</i> , <i>A. sclerotiiicarbonarius</i> , <i>A. sclerotioniger</i>
<i>A. heteromorphus</i> clade	<i>A. ellipticus</i> , <i>A. heteromorphus</i>
<i>A. homomorphus</i> clade	<i>A. homomorphus</i> , <i>A. saccharolyticus</i>
<i>A. aculeatus</i> clade	<i>A. aculeatinus</i> , <i>A. aculeatus</i> , <i>A. fijiensis</i> , <i>A. indologenus</i> , <i>A. japonicus</i> , <i>A. uvarum</i> , <i>A. violaceofuscus</i>

The taxonomy of section *Nigri* is one of the most confusing and complex in the fungal world due to the subtle differences between the species, which is underlined by several proposed taxonomic

schemes^[140,141,144,148,149]. As a result several reported strains have incorrectly been named *A. niger*, because the organism is seen as a black Aspergilli and therefore reported as *A. niger*^[146]. Using morphological structures can be a helpful tool when identifying black Aspergilli, especially a species, such as *A. carbonarius* (size of conidia) and the uniserate species (e.g. *A. japonicus*, *A. aculeatus*) are easily recognized using identification manuals^[144,145]. In contrast, species related to *A. niger* are very difficult to distinguish^[144]. A polyphasic identification approach, in which morphology, metabolite profiling, and/or several molecular methods (sequence data of the β -tubulin, calmodulin and ITS regions) are used, is thus preferred^[144,145,150]. An example of the problematic identification of species within section *Nigri* was given in a recent comparison of the β -tubulin, calmodulin and ITS sequence data from all described members of the section *Nigri*. This comparison showed that *A. foetidus* and *A. lacticoffeatus* are synonyms of *A. niger* and *A. coreanus* is synonym of *A. acidus*^[145].

1.3.2 The industrial workhorses

The black Aspergilli are able to produce a wide range of metabolites, as listed in section 1.3.3. Also, the black Aspergilli have a number of characteristics which make them ideal organisms for industrial applications as high levels of protein secretion and ability to assimilate various organic substrates^[151]. Therefore, many species of the black Aspergilli are used in the fermentation industry to produce hydrolytic enzymes, such as amylases or lipases, and organic acids such as citric acid and gluconic acid^[152,153]. The versatile metabolism of *A. niger* has made it one of the most important production organisms for industrial fermentation^[56,152], and after thorough research *A. niger* used with known industrial processes, has been placed on the Generally Recognized as Safe (GRAS) list of the USFDA^[154].

The first industrial use of *Aspergillus niger* was in 1919, when its ability to produce citric acid was exploited. *A. niger* is the preferred choice of microorganism for the citric acid bio-production process, due to a high ease of handling/harvesting, extraordinary ability to ferment a variety of raw materials (e.g. agricultural waste residues), and high product yields^[155]. Strain improvement and optimization of the production media have resulted in a process of citric acid accumulation by which cells convert up to 80% of the substrate, sucrose, to the final product^[156].

A. niger and *A. awamori* are well established expression platforms for the production of secreted proteins and enzymes^[157]. In the natural environment these fungi are involved in the degradation of plant cell material by producing highly specialized enzymes such as pectinases, cellulases, and xylanases^[151]. *A. niger* is additionally considered the ancestor of many currently used enzyme production strains^[56], and has been used as a rich source of a variety of enzymes well established in fruit processing, baking, and the starch and food industries since the 1960s. The first to be utilized were pectinase, protease, and amyloglucosidase^[152].

1.3.3 Secondary metabolism

The secondary metabolism of the black *Aspergilli* is wide and diverse, covering alkaloids, polyketide and terpenes. More than 145 metabolites have been isolated and structure elucidated from the biserate black *Aspergilli*^[158] a thorough review of this is given in paper 9. Examples of secondary metabolites produced by members of section *Nigri* are given in figure 3

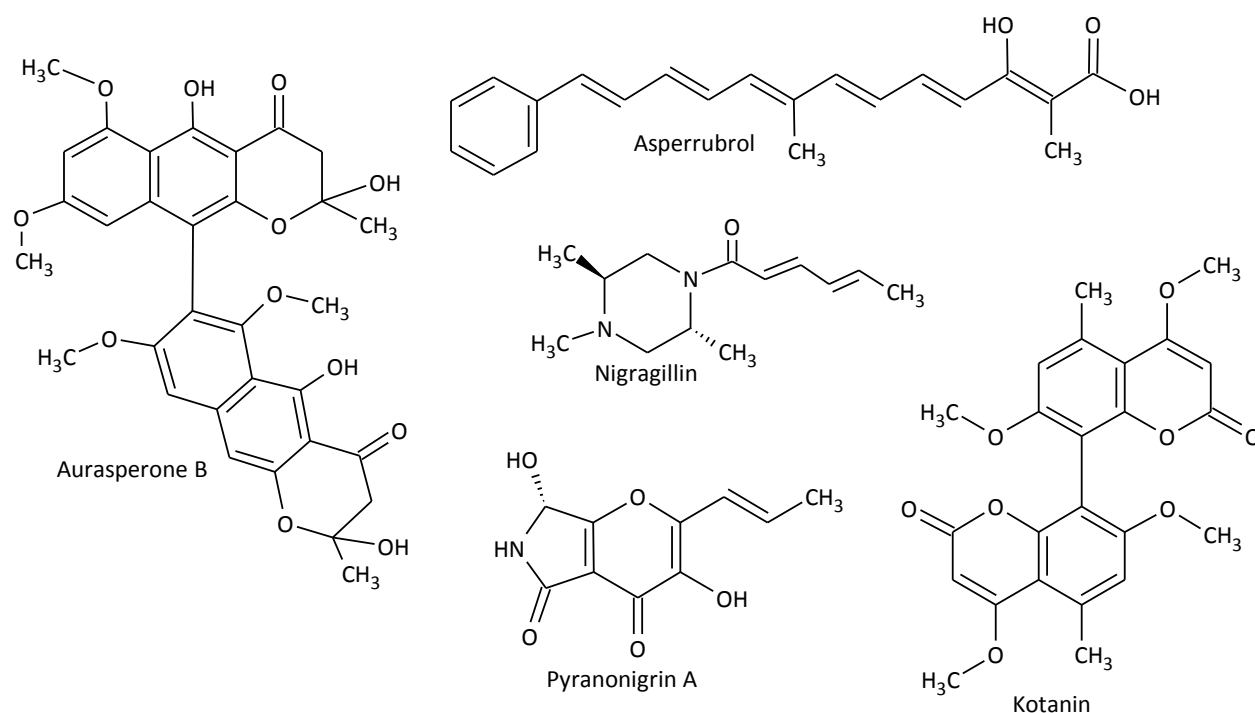


Figure 3: Examples of secondary metabolites produced within Section *Nigri*.

Several guidelines to distinguish between the species using the secondary metabolite profile can be used, since there are several differences among the species. Table 7 gives an overview of the most common secondary metabolites produced by black *Aspergilli*.

Table 7: Production of secondary metabolites in section *Nigri*^[139,144-146,159]

Species	Extrolite
<i>A. acidus</i>	antafumicins, asperazine, funalenone, naphtho- γ -pyrones, pyranonigrin A, nigragillin
<i>A. aculeatinus</i>	Aculeasins, neoxaline, secalononic acid D, secalononic acid F
<i>A. aculeatus</i>	secalononic acid D, secalononic acid F
<i>A. awamori</i> *	Same as <i>A. niger</i>
<i>A. brasiliensis</i>	naphtho- γ -pyrones, pyrophen, tensidol A & B
<i>A. carbonarius</i>	ochratoxins (A, B, α , β), naphtho- γ -pyrones, pyranonigrin A
<i>A. costaricensis</i>	aflavinines, funalenone, naphtho- γ -pyrones
<i>A. ellipticus</i>	austdiol, candidusins, terpenyllin, cf. xanthoascins
<i>A. eucalypticola</i>	Pyranonigrin A, funalenone, naphtho- γ -pyrones,
<i>A. fijiensis</i>	asperparalins, secalononic acid D, F
<i>A. heteromorphus</i>	lots of highly unique extrolites including indol-alkaloids, none of them structure elucidated
<i>A. homomorphus</i>	dehydrocarolic acid, secalononic acid D, secalononic acid F
<i>A. ibericus</i>	naphtho- γ -pyrones, pyranonigrin A
<i>A. indologenus</i>	okaramins A, B, H
<i>A. japonicus</i>	cycloclavine, festuclavine
<i>A. neoniger</i>	funalenone, naphtho- γ -pyrones, pyranonigrin A
<i>A. niger</i>	Fumonisin B ₂ , B ₄ , B ₆ , funalenone, kotanins, malformins, naphtho- γ -pyrones, nigragillin, ochratoxin A, pyranonigrin A, tensidol A & B
<i>A. piperis</i>	aflavinins, naphtho- γ -pyrones, pyranonigrin A
<i>A. saccharolyticus</i>	None known
<i>A. sclerotii carbonarius</i>	naphtho- γ -pyrones, pyranonigrin A, three unique indol-alkaloids at retention indices 1475, 1676 and 1838.
<i>A. sclerotium niger</i>	corymbiferan lactones, funalenone, naphtho- γ -pyrones, ochratoxins (A, B, α , β), pyranonigrin A
<i>A. tubingensis</i>	asperazine, funalenone, malformins, naphtho- γ -pyrones, nigragillin, pyranonigrin A, tensidol A & B
<i>A. uvarum</i>	asterric acid, dihydrogeodin, erdin, geodin, secalononic acid D and F
<i>A. vadensis</i>	asperazine, naphtho- γ -pyrones, nigragillin, a polar orlandin-like compound
<i>A. violaceofuscus</i>	indol-alkaloids

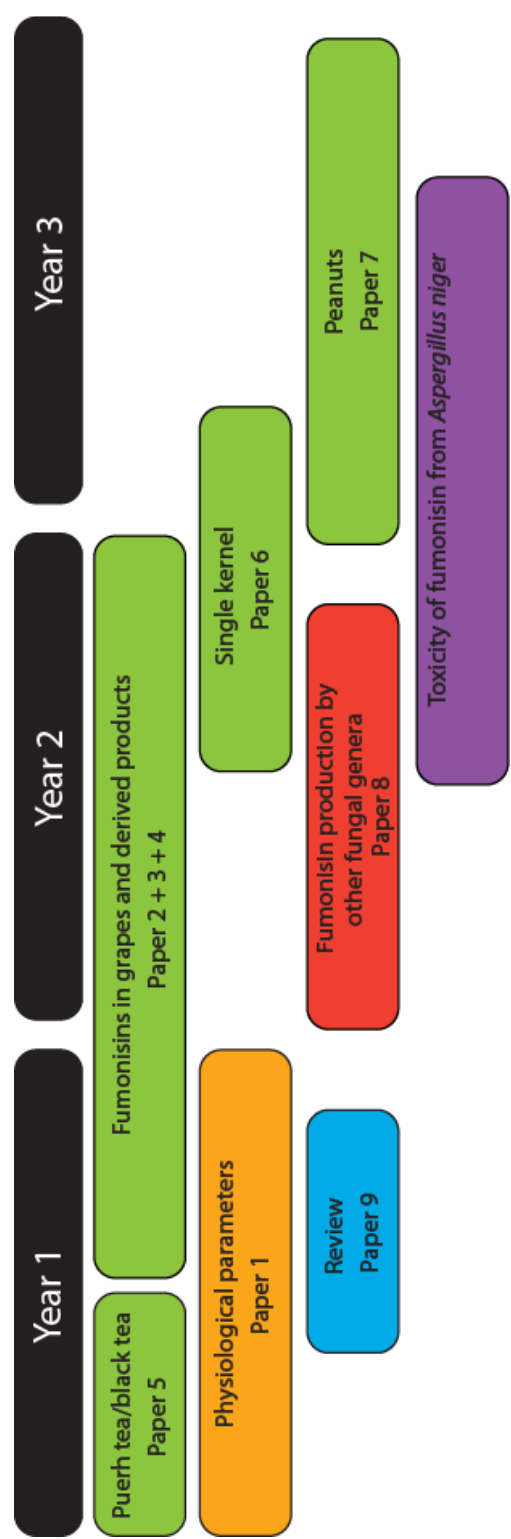
*: only a phylogenetic species

The economical most important secondary metabolite within section *Nigri* is ochratoxin A and its precursors due to its high prevalence in food and feed and toxicity. The main organisms producing this compound are *A. carbonarius* and *A. niger*; however, production of ochratoxin A by *A. japonicus*^[160,161], *A. sclerotium niger*^[139], and *A. tubingensis*^[162] has also been reported, although those of *A. japonicus* and *A. tubingensis* are questioned. *A. carbonarius* is seen as a consistent ochratoxin A producer, contrary to *A. niger* of which roughly 10-15 % produce this mycotoxin at 10-1000 lower amounts^[146,163,164]. Another major group of mycotoxins, produced within section

Nigri is the fumonisins; however, so far only *A. niger* is reported as a producer of this compound. Other species tested with a negative result are *A. acidus*, *A. aculeatinus*, *A. aculeatus*, *A. brasiliensis*, *A. carbonarius*, *A. costaricaensis*, *A. ellipticus*, *A. heteromorphus*, *A. homomorphus*, *A. ibericus*, *A. japonicus*, *A. piperis*, *A. sclerotii carbonarius*, *A. sclerotium niger*, *A. tubingensis*, *A. uvarum*, and *A. vadensis*^[137,146,165]. This means there are still 6 species that have not been screened for a fumonisin production, all newly described^[145]. Several reports indicate that 60-80% of *A. niger* strains are able to produce fumonisins^[137,165], indicating that the fumonisin production by *A. niger* is more significant than OTA from *A. niger*. Since *A. niger* are able to produce both mycotoxins, a simultaneously production can occur, and approximately 10% are able to produce both mycotoxins^[146]

2 Overview of experimental work

This figure is intended as an overview of the main experiments performed during the period. The obtained results will be discussed in the following chapters. The experimental design is found in the denoted papers.



3 Results and Discussion

3.1 Comparative physiological studies of fumonisin production in *Fusarium* and *Aspergillus niger*

When this project was initiated, the knowledge and understanding of the fumonisin production by *A. niger* was sparse. It was only known that *A. niger* produced FB₂ preferably on media with a high sugar/salt content^[34]. Therefore, it seemed relevant to investigate how physiological parameters like temperature and water activity affected fumonisin production by *A. niger*.

To compare *A. niger* and *Fusarium*, 25 *Fusarium* strains and five *A. niger* strains were inoculated on three different media, Czapek yeast extract agar + 5% NaCl (CYAS)^[166], Rice meal corn steep liquor agar (RC)^[167], potato dextrose agar (PDA)^[147] and incubated for seven days at 25 °C. The *A. niger* strains consistently produced FB₂ on all three media favoring RC slightly over CYAS and with a poor production on PDA, table 8. These results partly agree with what Frisvad *et al.*^[34] stated, CYAS only supported fumonisin production by *A. niger* and RC supports fumonisin production by both genera. Fumonisin production was detected in all five *A. niger* strains on PDA, which is contradictory to the study by Frisvad *et al.* where no fumonisin production was detected, though they only tested one strain and the lack of fumonisin production could be explained by levels were below the detection limit or strain differences, as our results indicate occurs in *Fusarium* species.

Table 8: Fumonisin production by *Aspergillus niger* (FB₂) and *Fusarium* spp. (FB₁, FB₂, FB₃) on CYAS, PDA and RC after 7 days growth at 25°C. The replicates are made in biological duplicates on two separate plates.

Fungi	CYAS µg/ml	RC µg/ml	PDA µg/ml
<i>A. niger</i> (n=5)	2.9-25 (5/5)	5.3-35 (5/5)	0.46-3.1 (5/5)
<i>F. proliferatum</i> (n=6)	n.d. (0/6)	0.028-46 (6/6)	2-33 (4/6)
<i>F. verticillioides</i> (n=6)	n.d.(0/6)	0.028-2.2 (6/6)	4.5-35 (5/6)
<i>F. dlamini</i> (n=2)	n.d. (0/2)	n.d. (0/2)	n.d. (0/2)
<i>F. napiforme</i> (n=2)	n.d. (0/2)	0.13-0.24 (2/2)	0.081-6.2 (2/2)
<i>F. nygamai</i> (n=8)	n.d. (0/8)	0.01-22(8/8)	0.14-16 (6/8)
<i>F. oxysporum</i> (n=1)	n.d. (0/1)	2.78 (1/1)	37 (1/1)

CYAS: Czapek Yeast extract Agar + 5%NaCl; PDA: Potato Dextrose Agar; RC: Rice meal corn steep liquor agar.

Standard deviation calculated on two measurements

n.d. not detected.

For further analysis, the five *A. niger* and one randomly selected strain from each *Fusarium* species were inoculated on CYAS and PDA respectively and incubated at six different temperatures between 15-42 °C for 7 days. Fumonisin production occurred at temperatures between 20-37 °C

(figure 4), with four strains having a maximum production at 25 °C and the latter at 30 °C. At 42 °C all *A. niger* strains grew, but had no detectable fumonisin production. Thus production of fumonisin seem to follow a similar pattern to that of ochratoxin A^[168]. However on a synthetic grape medium and maize kernels ochratoxin production were largest at 15 °C^[169,170].

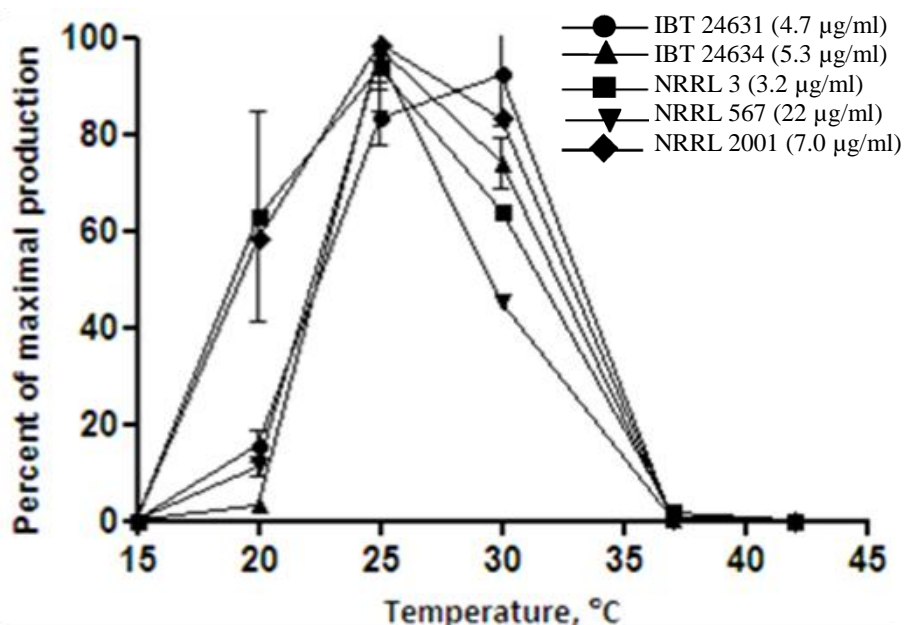


Figure 4: FB₂ production by *A. niger* (A) after 7 days of growth at different temperatures (15-42°C) on CYAS. The concentration of FB₂ was detected in the methanol:water (3:1) extract. The values are means of biological duplicates on two different plates; highest value (µg/ml) is in parenthesis. (PAPER 1)

Fumonisin production at different a_w by two of the five strains (IBT 24634 and NRRL 567) is given in figure 5. All five strains had decreasing fumonisin production, with decreasing water activity, when grown on glycerol. On sucrose both had an increase in their production; however, below 0.99 IBT 24634 had decreasing production, ending below the production at starting conditions. NRRL 567 had more or less constant production below 0.995. On NaCl a different picture was seen, NRRL 567 had a slight increase followed by a decreasing production. IBT 24634 had a relatively constant production, with no variations until 0.94 after which it stops. In agreement with our results, Frisvad *et al.* found that the addition of 5 % NaCl ($a_w = 0.97$) or 20 % sucrose ($a_w = 0.99$) increased the production of FB₂ by *A. niger*^[34].

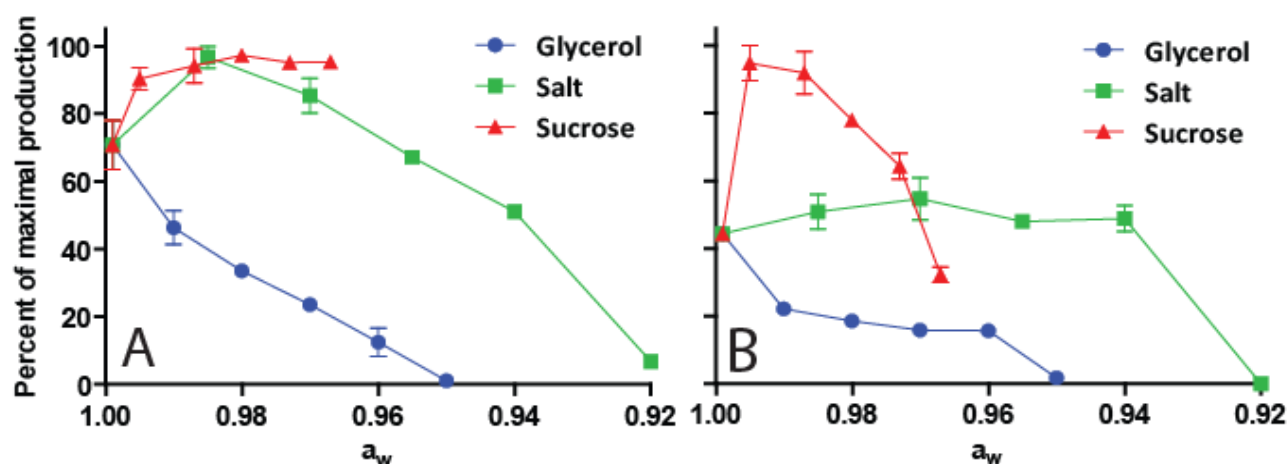


Figure 5: Effect of glycerol, NaCl and sucrose on the FB_2 production of *A. niger*, A: NRRL 567 (max: 22 $\mu\text{g/ml}$) B: IBT 24634 (max: 5.3 $\mu\text{g/ml}$). Isolates have been incubated at 25 °C for 7 days on CYA. The concentration of FB_2 was detected in methanol:water (3:1) extracts. The values are means of biological duplicates on two different plates. (PAPER 1)

As this is currently the only report of the effect of water activity on the production of fumonisins by *A. niger*, a comparison is made to the production of ochratoxin A by *A. niger*. Previous studies of the optimal water activity of the production of ochratoxin A are inconclusive, stating both 0.95 and the range 0.96-0.99^[170,171]. These are different from our results, with four strain preferring 0.99 and the last 0.98. Each strain also responded differently to the various solutes, with highest production on sucrose, followed by NaCl and glycerol.

Earlier studies of the optimal a_w for fumonisin production by *Fusarium*, showed that it favored 0.97-0.98^[172,173]. Our results furthermore indicate, that the regulation of the fumonisin production in *A. niger* and *Fusarium* are very different, with the latter preferring high a_w (>0.99) and lower temperatures (20-25 °C), compared to *A. niger* which prefers lower a_w and higher temperature (25-30 °C).

3.2 Fumonisin from *Aspergillus niger* in food

3.2.1 Production of fumonisin in grapes and derived products

Due to the presence of ochratoxin A in grapes and derived products, black Aspergilli have already been identified as the causative organisms in these commodities^[174-176]. The fumonisin contamination of grapes and their derived products was obviously relevant to investigate. In order to determine the distribution of fumonisins producing *A. niger* in raisins, black Aspergilli were isolated from 17 different raisin brands. Altogether 86 black Aspergilli were isolated and 66 were

identified as *A. niger*, 4 *A. tubingensis*, and 16 *A. acidus*. These isolates were screened for potential fumonisin production on three different media, Czapek yeast extract agar (CYA)^[166], yeast extract sucrose agar (YES)^[166] and CYAS^[166]. Neither *A. tubingensis* nor the *A. acidus*, produced detectable amounts of fumonisins, as opposed to 77 % of the *A. niger* strains, that were able to produce fumonisins. This frequency was similar to that reported for *A. niger* from green coffee beans^[36] and grapes^[137], as well as industrial *A. niger* strains^[146]. Though higher than those reported from Californian raisins^[165] and Portuguese grapes^[177]. The strains isolated in the Portuguese grape study were only identified as *A. niger* aggregate, which explains the lower frequency, since this also covers other common black Aspergilli like *A. tubingensis* and *A. acidus*. The abundance of fumonisin producing *A. niger* confirmed that a potential fumonisin contamination was present. Worst case scenarios were performed by inoculation of ten isolates on grapes. All ten isolates were able to produce fumonisins on the grapes, table 9. Surprisingly none of the ten isolates were able to produce ochratoxin A under these circumstances.

Table 9: Production of fumonisins by *Aspergillus niger* in grapes after 7 days growth at 25°C. (PAPER 2)

Isolate IBT	Fumonisin B ₂ µg/kg		Fumonisin B ₄ µg/kg ^A		Fumonisin B ₆ µg/kg ^B		Ochratoxin A µg/kg ^C
	Sample 1	Sample 2	Sample 1	Sample 2	Sample 1	Sample 2	
28747	951	1283	75	189	n.d.	n.d.	n.d.
28753	7703	7979	772	1542	117	129	n.d.
28934	2098	5376	158	612	5	77	n.d.
28937	1045	1941	253	265	n.d.	n.d.	n.d.
28948	3380	4772	349	509	63	65	n.d.
28964	91	251	5	23	n.d.	n.d.	n.d.
28965	1074	2414	148	344	28	n.d.	n.d.
28966	1261	5123	220	634	60	n.d.	n.d.
28994	432	2422	36	158	n.d.	n.d.	n.d.
29019	4303	11189	143	405	89	135	n.d.

^A: Assuming same response factor as fumonisin B₂. ^B: Assuming same response factor as fumonisin B₁. ^C: LOD: 3 µg/kg. nd: not detected, LOD: 5 µg/kg

All strains produced FB₂ and FB₄, whereas only six strains produced FB₆, the average ratio of these is ~89% FB₂, ~10 % FB₄ and ~1 % FB₆. The FB₂ levels in grapes were 171-7841 µg/kg, FB₄ were 14-1157 µg/kg, and FB₆ levels were 5-135 µg/kg. These fumonisin levels are comparable to a similar study by Logrieco *et al.*^[137].

To investigate how/if the fumonisin production could occur in raisins, four of the isolates were selected for further analysis. To imitate real life scenarios, dried grapes and two types of re-moistened raisins were used. The dried grapes had three different initial water activities (0.54;

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0.69; 0.76). The mean FB₂ and FB₄ production in the dried grapes were 91-1747 µg/kg and 9-69 µg/kg, table 10. Fumonisin B₆ was only found in two samples at very low levels. All samples had lower fumonisin levels compared to fresh grapes. The overall trend from this were, that the fumonisin production decreased with decreasing a_w, which correlates with our earlier results, that lower a_w results in a low fumonisin production (Chapter 3.1).

Table 10: Production of fumonisins by *A. niger* in dried grapes after 7 days growth at 25°C.^A(PAPER 2)

Isolate	Weight loss %	Initial a _w	Fumonisin B ₂ µg/kg	Fumonisin B ₄ µg/kg ^B	Fumonisin B ₆ µg/kg ^C	Ochratoxin A µg/kg ^D
28753	32	0.76±0.04	1747±316	48±22	n.d.	n.d.
	38	0.69±0.01	1128±329	48±16	8±2	n.d.
	52	0.54±0.09	91±73	9±4	n.d.	n.d.
28934	32	0.76±0.04	1363±954	69±51	7±5	n.d.
	38	0.69±0.01	1033±460	50±23	n.d.	n.d.
	52	0.54±0.09	636±397	24±16	n.d.	n.d.
28948	32	0.76±0.04	313±323	54±47	n.d.	n.d.
	38	0.69±0.01	154±71	27±11	n.d.	n.d.
	52	0.54±0.09	144±104	29±19	n.d.	n.d.
29019	32	0.76±0.04	203±106	25±11	n.d.	n.d.
	38	0.69±0.01	214±137	28±13	n.d.	n.d.
	52	0.54±0.09	102±76	10±5	n.d.	n.d.

^A: The weight loss was 32% (4 h), 38% (5 h), and 52% (6.5 h). The values are means of the quadruplicates, plus/minus the standard deviation, LOD: 5 µg/kg. ^B: Assuming same response factor as fumonisin B₂. ^C: Assuming same response factor as fumonisin B₁. ^D: LOD: 3 µg/kg. nd: not detected.

Two types of re-moistened raisins were made each with different initial a_w (0.77;0.95), both types initially surface sterilized. **Scenario 1**: Raisins were placed on a filter placed on water agar, allowing the raisin to be kept at stagnate or increasing a_w, figure 6. **Scenario 2**: Raisins were covered with water and kept overnight at 5 °C and placed in an empty Petri dish, the raisin were hereby kept in an environment with decreasing a_w, figure 6.

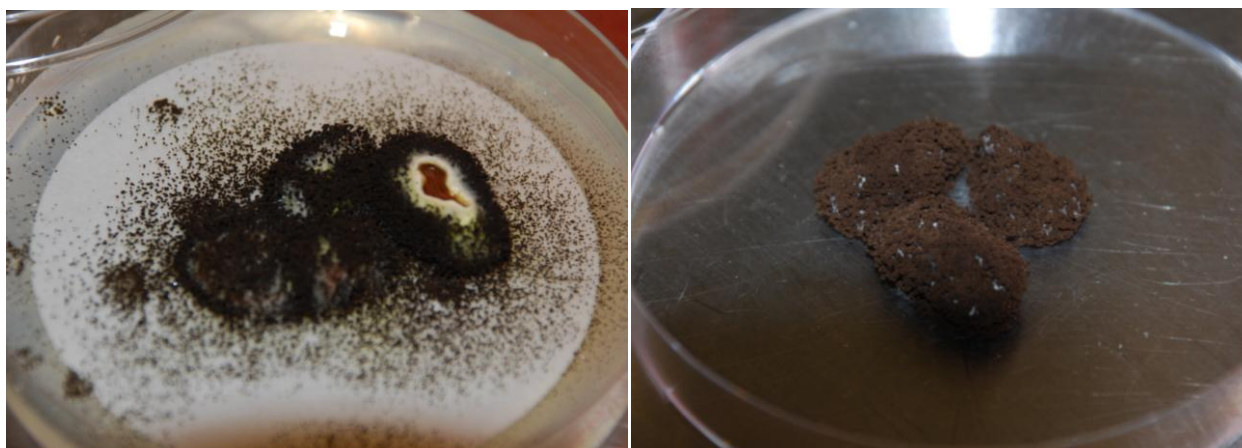


Figure 6: Two types of worst-case scenarios of re-moistened raisins. Left: Re-moistened raisins were placed on a filter on top of the water agar. Right: Raisins were covered with water and kept overnight at 5 °C and placed in an empty Petri dish

Fumonisin production in all samples was markedly higher in scenario 1 (229-6476 µg/kg) than scenario 2 (5-784 µg/kg), table 11. Only two strains produced FB₆ (9-68 µg/kg) at low levels compared to FB₂ and FB₄.

Table 11: Production of fumonisins by *A. niger* in grapes after 7 days growth at 25 °C^A. (PAPER 2)

Isolate IBT	Weight gain	Initial a _w	Fumonisin B ₂ µg/kg	Fumonisin B ₄ µg/kg ^B	Fumonisin B ₆ µg/kg ^C	Ochratoxin A µg/kg ^D
28753	7% (agar)	0.77±0.05	1160±686	105±56	22±12	n.d.
	50% (water)	0.95±0.001	407±249	47±25	7±5	n.d.
28934	7% (agar)	0.77±0.05	229±58	27±7	n.d.	n.d.
	50% (water)	0.95±0.001	5±5	n.d.	n.d.	n.d.
28948	7% (agar)	0.77±0.05	459±430	45±31	n.d.	n.d.
	50% (water)	0.95±0.001	112±51	12±4	n.d.	n.d.
29019	7% (agar)	0.77±0.05	6476±1139	356±46	68±12	n.d.
	50% (water)	0.95±0.001	784±636	672±422	9±5	n.d.

^A: The values are means of the quadruplicates, plus/minus the standard deviation, LOD: 5 µg/kg.

^B: Assuming same response factor as fumonisin B₂.

^C: Assuming same response factor as fumonisin B₁.

^D: LOD: 3 µg/kg. nd: not detected

Since *A. niger* is able to produce fumonisins in grapes, the derived products needed to be looked into, especially because of the previously reported problems with ochratoxin A in wine and raisins^[28,174,178-180], as well as the findings of FB₂ in grape must^[137]. As a result, it was decided to investigate retail raisins. Totally, 21 raisin brands were screened for fumonisins. FB₂ and FB₄ were detected in 10 samples. The detected levels were FB₂: 1.3 -13 µg/kg and FB₄: 0.31-1.3 µg/kg, table 12. Of the 10 positive samples 7 originated from California, 2 from Chile and 1 from Turkey.

Table 12: Overview of the positive raisin samples and their average fumonisin content^a. (PAPER 4)

Brand	Origin	Grape variety	Organic	Fumonisin B ₂ (µg/kg)	Fumonisin B ₄ (µg/kg) ^b
1	California	Thompsons Seedless		4.9±0.4	0.26±0.05
				4.5±1	0.8±0.4
				13.0±1	1.3±0.4
				7.5±0.8	0.9±0.2
2	California	Thompsons Seedless		2.7±0.2	0.29±0.1
3	California	Thompsons Seedless	✓	1.8±0.2	<LOQ ^c
4	California	Thompsons Seedless	✓	2.6±0.4	0.83±0.5
5	California	Thompsons Seedless		3.8±0.6	0.31±0.09
6	California	Thompsons Seedless		4.1±0.6	1.0±0.1
7	California	Thompsons Seedless	✓	1.3±0.6	0.25±0.2
12	Turkey	Sultana		4.8±2	0.75±0.1
20	Chile	Golden		<LOQ	ND
21	Chile	Golden		1.4±0.6	<LOQ

^a Fumonisin concentration is given as the average of three determinations from the same package, ±SD.

^b Fumonisin B₄ was calculated assuming a 4 fold better response factor than for fumonisins B₂ shown at one spiking experiment. Fumonsin B₁ and B₆ were not detected in any of the packages.

^c LOQ, limit of quantification: fumonisin B₂: 1 µg/kg and for B₄ estimated to 0.25 µg/kg.

^c ND, not detected, limit of detection for fumonisin B₂ was 0.3 µg/kg and for B₄ 0.1 µg/kg

To further investigate the magnitude of the fumonisin contamination in grape derived products, we chose to look at wine, and since there, at that point, was no method for detection of fumonisins in wine, a prime objective for this project was development of a method. For this purpose several SPE columns were tried. Initially, SAX was the choice, however irreproducibility and poor recovery urged for a different approach. Instead cation-exchange was attempted, targeting the amine group of the B-series fumonisin. Testing of six different cartridges (mixed-mode and polymeric), Strata-X-C and OASIS MCX was found to be superior to the others. Due to a slightly better reproducibility Strata X-C were chosen. The method was validated showing a mean recovery of 83% with an RSD of 12%. Using this method, 77 wine samples were screened for fumonisin content, and 23% of these were positive (n=18), with fumonisin levels between 1-25 µg/L, table 13. The frequency and levels were similar to a contemporary study of Italian wines^[114], which showed a positive fraction of 17% and fumonisin levels between 0.4-2.4 µg/L.

Table 13: Overview of the positive wine samples^a. (PAPER 3)

Sample number	Country	Grape sort	Winetype	Year	Fumonisin B ₂ (µg/L)
10	Australia	Shiraz	Red	2008	1.0
24	France	Carignan, Grenache, Syrah	Red	2003	5.1
37	France	Grenache, Syrah	Red	2003	1.3
68	France	Grenache, Syrah, Mourvedre	Red	2008	6.5
9	Italy	Merlot	Red	2007	7.0
14	Italy	Corvina, Rondinella, Molinarea, Negrana	Red	2007	6.7
39	Italy	Montepulciano, Aglianico	Red	2005	2.1
40	Italy	Merlot, Corbina	Red	2007	3.9
58	Italy	Negroamaro, Primitivo	Red	2006	1.6
63	Italy	n.s.	Red	2007	1.2
66	Italy	Corvina, Rondinello, Molinara	Red	2004	3.2
89	Italy	n.s.	White	2008	1.0
104	Portugal	n.s.	Port	n.s.	2.8
7	Romania	Pinot Noir	Red	2006	20
5	South Africa	Cabernet Sauvignon	Red	2007	2.4
2	Spain	Monastrell	Red	n.s.	2.1
27	Spain	Tempramito, Gaunacha	Red	2005	1.9
33	USA, CA	Zinfandel	Red	1998	25

^aThe description of the wines was read off the label of the bottle

n.s.: Not specified

Of the 18 positive samples, 16 were red, one white and one port wine. A similar high frequency of mycotoxin in red wine is also seen with ochratoxin A contamination, which is probably caused by the differences in winemaking^[175,178,181]. Furthermore, results indicated, although not significantly, that Italian wine (35%) was more frequently contaminated compared to Spanish (22%) and French (14%) wine, although at lower average levels than French but higher than the Spanish.

The levels found in the raisins (1.4-14.3 µg/kg) and wine (1-25 µg/L) as well as those found in coffee beans^[36] (1-9.7 µg/kg) are comparable; however, the frequency of contaminated samples are more than twice as large in raisins and coffee beans than wine (23 %) with 48 % and 58 %, respectively. On the basis of the tested wine and raisin samples, it can be concluded that the frequency of heavily infected grapes is low, since the detected levels are 100-400 times lower than those found in highly contaminated grapes in this study, and a prior by Logrieco *et al.*^[137]. This could be due to efficient removal of infected/damaged berries, initiated after problems with ochratoxin A in grapes and derived products were reported in the late 1990s^[178,182,183].

The detection of FB₄ in retail raisins agree with the previous results from artificially infected grapes and raisins; however, fumonisin B₄ was not identified in the studies of fumonisins in wine^[114] and grape must^[137]. During raisin production there is the possibility of both pre- and post-harvest contamination, since the grapes are often sun-dried^[184]. The reason for the FB₄ production, could

be caused by a different growth physiology and metabolism of *A. niger* during the drying process of the grapes. Which is probably caused by a high growth rate and as a consequence, some of the FB₄ is not converted into FB₂, before the water activity decreases below *A. niger* growth limits (~ 0.85)^[185]. The postharvest growth together with the water loss during the drying process of the vine fruits and the simultaneous concentration of the fumonisins, may explain why fumonisin B₄ is not detected in this, and other wine studies^[137].

3.2.2 Fumonisin from *Aspergillus niger* in other food commodities

Besides grape, wine, and raisins where *A. niger* is a considered huge problem, other commodities where *A. niger* could pose a risk, were also looked at. Black tea/puerh tea and peanut/peanut butter were chosen. The puerh tea was chosen because it was reportedly fermented with *A. niger*, which also is the dominant organism in black tea^[186-190]. Ten teas were investigated, five regular black tea and five puerh tea. Incubation of the teas on DG18 led to the isolation of 47 black *Aspergilli*. To determine their metabolite profile including the potential fumonisin production, the strains were analyzed by LC-MS and HPLC-UV. None of the 47 strains had detectable amounts of fumonisins or ochratoxin A, but other metabolites were identified. All strains produced asperazine with 12 furthermore producing antafumicin, consequently none of the strains could be *A. niger*, since these compounds are not associated with *A. niger*^[144]. Production of antafumicin is only found in *A. acidus*, and thus the 12 strains could be characterized as this species. From their metabolite profile the remaining 35 strains were provisionally identified as *A. acidus* or *A. tubingensis*. Sequences of a part of the calmodulin gene in 17 of these strains, were thus examined. The partial sequences were identical to that of *A. acidus*, and all 35 strains were hereby identified as *A. acidus*. Since all black *Aspergilli* were identified as *A. acidus*, no obvious risk of mycotoxin contamination would be associated with black and puerh tea. The suggested non-toxicity is supported by a toxicity study, in which no adverse effects, were observed after intake of puerh tea extracts^[191].

An additional food commodity was investigated; this was peanut and peanut butter. The potential for fumonisin production in peanuts were determined by worst case scenarios by inoculation of *A. niger* on shelled peanuts, figure 7.



Figure 7: Examples of worst-case scenarios of *A. niger* contaminated peanuts after 7 days growth at 25°C

Four *A. niger* strains were inoculated on peanuts and incubated for 7-14 days at 25 °C. The mean production of FB₂ varied more than 40 fold from 0.22 mg/kg (IBT 29019) to 9.4 mg/kg (IBT 18741). The measured fumonisin concentrations (table 14) are similar and comparable to those previously detected in grapes and raisins (0.2-8 mg/kg)^[137] (Chapter 3.2.1).

Table 14: Production of fumonisins by *Aspergillus niger* in peanuts after 7-14 days growth at 25 °C

Incubation period	IBT 18741 (FB ₂ , mg/kg)		IBT 28753 (FB ₂ , mg/kg)		IBT 29019 (FB ₂ , mg/kg)		IBT 29331 (FB ₂ , mg/kg)	
	#1	#2	#1	#2	#1	#2	#1	#2
7 days	5.8	4.8	3.5	1.6	0.43	0.014	2.5	1.0
14 days	9.6	9.1	2.5	3.9	0.56	3.4	3.9	7.6

Since *A. niger* is able to produce fumonisins when growing on peanuts, the potential for fumonisin contamination was present and peanut retail products were therefore screened. Analysis of 15 peanut samples (from local markets in Thailand) and nine peanut butter brands (peanuts originating from four countries) all analyzed in triplicate showed no indications of fumonisins (LOD: 0.004 mg/kg). Despite *A. niger* is a common contaminant of peanuts^[143,192-196], and that when grown on the peanuts, fumonisins are produced in high quantities (up to 2000 higher than our detection limit). We assume that the combination of: i) extensive damage to the nuts made

by *A. niger* (in some cases they are heavily digested), and: ii) the high attention to reduce aflatoxin exposure^[24] effectively removes *A. niger* infected peanuts and thus fumonisins.

3.3 Single kernel analysis of fumonisins from *Fusarium* in maize

The main crop where fumonisins are a concern is maize. Several reports from around the world, have described fumonisin contamination of maize and derived products. A new approach was initiated within this thoroughly investigated area, by investigating single kernels, instead of the traditionally approach, where the average content in lots was determined. Using this approach, the aim was to clarify, the effect of manual visual sorting, as well as the content and distribution of fumonisins in the single kernels. The maize was collected from subsistence farmers in South Africa, who had classified the maize in two categories: “Good” and “Moldy”. In total ten batches were investigated, five from each category.

The infection rate of the ten batches was determined by random selection of 100 kernels and counting of infected/damaged kernels. Good quality batches had infection rates below 3%, while the moldy batches had a marked difference, with infection rates of 6-19%, table 15, p29. To survey the mycobiota of the maize, visibly infected/damaged kernels were placed on four different media (24 per batch, $n_{\text{total}}=240$) and strains were isolated after 7 days growth at 25 °C. *Fusarium* infections were seen in 17-83 % of the kernels by primarily *F. verticillioides* and *F. subglutinans*. These findings were in agreement with previous studies^[197,198]. Other species found were *Penicillium aurantiogriseum*, *P. brevicompactum*, *P. concavorugulosum*, *P. crustosum*, *P. pittii*, and *Aspergillus wentii*.

Analysis by LC-MS/MS of infected/damaged (n=300) and uninfected/undamaged (n=100) kernels (figure 8) for fumonisin contamination, showed that 15% had detectable amounts of fumonisins and 4% were considered highly infected, because the fumonisin levels were above 100 mg/kg. None of the visually uninfected/undamaged kernels contained fumonisins. However, in the infected/damaged kernels fumonisin levels were batch dependent and varied from 3-40% with a total fumonisins (FB₁, FB₂, FB₃, and FB₄) range of 1.8-1428 mg/kg. Even though low-quality kernels were more frequently found in the poor-quality maize, low quality kernels from the good-quality maize had generally higher fumonisin content than those from the moldy batches.



Figure 8: Infected/damaged (A) and uninfected/undamaged (B) kernels

The fumonisin content in the batches were estimated using infection rate, average kernel weight, batch weight, and average fumonisin content. Good quality batches had an average estimated fumonisin content of 0.28-1.1 mg/kg and moldy batches contained 0.03-6.2 mg/kg. Two of the batches exceeded the maximum guidance level of 4 mg/kg in maize for human consumption recommended by the USFDA^[21] and four exceeded the European union regulatory limit of 1 mg/kg^[37].

Since only 15 % of the 400 kernels were positive for fumonisins, 80 samples were send to Austria for a multi-mycotoxin screen^[199] to identify toxins from the non-fumonisin producers. The 80 samples were selected to represent a combination of high, low, and no amount of fumonisins. Overall, 21 compounds were detected in the 80 kernels; however four kernels did not contain any detectable amounts of the analyzed toxins. Fumonisin were detected in 50 samples, of which one was characterized as uninfected, table 16, p30. Other frequently detected compounds were fusaric acid (n=53), emodin (n=27), chanoclavine (n=25), equisetin (n=14), and fusaproliferin (n=12). Seldom found compounds were agroclavine (n=3), altertoxin-I (n=3), beauvericin (n=7), moniliformin (n=4), nivalenol (n=4), rugulosin (n=4), and tentoxin (n=3). Alternariol, calphostin C, and cyclosporins A, C and H, were also detected, however only in single cases.

Table 15: Infection rate of baches and fumonisin (FB) content in uninfected and infected maize kernels (n_{total}=400).

Batch	416	438	440	447	448	404	429	431	432	455
Quality ^a	Good	Good	Good	Good	Good	Mouldy	Mouldy	Mouldy	Mouldy	Mouldy
Infection rate (%) ^{b,c}	66	66	83	66	100	100	83	66	83	83
Dominating FB-producing fungi (%) ^{b,d}	<i>F.sub</i> (17)	<i>F.vert</i> (17), <i>F.sub</i> (17)	<i>F.vert</i> (17)	<i>F.vert</i> (50)	<i>F.sub</i> (33), <i>F.vert</i> (17)	<i>F.vert</i> (50), <i>F.sub</i> (33)	<i>F.vert</i> (17)	<i>F.vert</i> (17)	<i>F.sub</i> (33)	<i>F.vert</i> (17)
Other fungi ^{b,c,d}	<i>P.conca</i> , <i>A.wentii</i> , <i>P.auran</i>	<i>P.conca</i> , <i>P.auran</i>	<i>P.conca</i>	<i>A.wentii</i>		<i>Eurotium</i> sp.		<i>P.crust</i> , <i>A.wentii</i>	<i>P.pittii</i> , <i>P.brevi</i>	<i>P.conca</i> , <i>P.crust</i>
Visibly infected or damaged (n=100)	2%	3%	3%	3%	3%	6%	14%	7%	19%	12%
Uninfected kernels contaminated with FB (n=10)	0%	0%	0%	0%	0%	0%	0%	0%	0%	0%
Visually infected kernels contaminated with FB (n=30)	20%	27%	27%	13%	10%	23%	10%	40%	23%	3%
FB contents in infected kernels (min-max, mg/kg)	34.4-1428	4.0-90	1.9-715	1.8-432	5.8-929	2.1-257	39-736	3.2-968	4.6-713	8.37
Mean FB content in infected kernels (mg/kg) ^e	330	49	114	149	328	54.8	297	166	159	8.37
Median FB in infected kernels (mg/kg)	53	42	30	82	50	4.8	117	33	41	8.4
Total FB content (mg/kg) ^f	1.1	0.28	0.79	0.52	0.85	0.67	3.7	4.1	6.2	0.03

Note: Fumonisin content is the total FB₁, FB₂, FB₃ and FB₄ content.Abbreviations: *F.vert*: *F. verticillioides*; *F.sub*: *F. subglutinans*; *P.auran*: *P. aurantiogriseum*; *P.brevi*: *P. brevicompactum*; *P.conca*: *P. concavurugulosum*; *P.crust*: *P. crustosum*.^a As sorted by the five subsistence farmers.^b Surface sterilization in hypochlorite prior to plating for a total of 180 randomly selected kernels (18 per batch with 6 on DG18%, 6 on CZID, and 6 on PCMA).^c Plating of 60 visibly infected or damaged kernels on water agar per batch.^d Not including sterile mycelia and species found only once among the 240 kernels.^e LC-MS/MS on Ultima QQQ. LOD_{FB1} of 0.16 mg/kg and LOD_{FB2} of 0.057 mg/kg^f Total FB content were calculated from the infection rate, mean kernel weight, batch weight, and mean fumonisin concentration.

Table 16: Distribution, mean and likely producer of metabolites found in visual infected and uninfected kernels.

Mycotoxin	% in uninfected kernels (n=8)	Mean in uninfected kernels	% in infected kernels (n=72)	Mean in infected kernels	Likely producer	Supported by microbiology	Produced pre or post harvest
Alternariol	0	-	1	13.8 µg/kg	<i>Alternaria</i>	No	Pre ^a
Altartoxin-I	13	0.26 µg/kg	3	0.18 µg/kg	<i>Alternaria</i>	No	Pre ^a
Beauvericin	13	0.32 µg/kg	8	8.4 µg/kg	<i>F. subglutinans</i>	Yes	Pre
Calphostin C	0	-	1	58.1 µg/kg	<i>Cladosporium</i>	No	Pre ^a
Cyclosporins	13	6.5 mg/kg	0	-	<i>F. solani</i> or <i>Tolypocladium</i>	No	Pre
Equisetin	25	83 µg/kg	17	75 µg/kg	<i>F. equiseti</i>	No	Pre
Fumonisin B ₁	0	-	60	27 mg/kg	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fumonisin B ₂	13	0.0035 mg/kg	65	8.7 mg/kg	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fumonisin B ₃	0	-	51	2.1 mg/kg	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fumonisin B ₄	0	-	49	1.6 mg/kg	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fusaproliferin	0	-	17	11 µg/kg	<i>F. subglutinans</i>	Yes	Pre
Fusaric acid	13	0.2mg/kg	64	34 mg/kg	<i>F. verticillioides</i> , <i>F. subglutinans</i> and other related	Yes	Pre
Moniliformin	0	-	6	202 µg/kg	<i>F. subglutinans</i>	Yes	Pre
Nivalenol	13	2.01 µg/kg	4	59 µg/kg	<i>F. equiseti</i> , <i>F. graminearum</i>	(Yes) ^b	Pre
Tentoxin	13	0.37 µg/kg	3	0.12 µg/kg	<i>Alternaria</i>	No	Pre ^a
Agroclavin	13	0.13 µg/kg	3	0.25 µg/kg	<i>P. concavurugulosum</i>	Yes	Post
Chanoclavine	38	4.5 µg/kg	31	1.2 µg/kg	<i>P. concavurugulosum</i>	Yes	Post
Emodin	75	0.73 µg/kg	29	2.2 µg/kg	<i>Eurotium</i> , <i>A. wentii</i>	Yes	Post
Rugulosin	0	-	6	372 µg/kg	<i>P. concavurugulosum</i>	Yes	Post

The majority of these compounds are presumably produced pre-harvest derived from the naturally growth of *Alternaria* and *Fusarium* in the field. The remaining part, is assumed to be produced post-harvest, since these are produced by *Penicillium*, *Aspergillus*, and *Eurotium*. Kernels with fusaproliferin and/or moniliformin had often lower amounts of fumonisins, indicating the presence of *F. subglutinans*, which is known as a poor fumonisin producer^[200]. Additionally, kernels with both fusaric acid and fumonisins had an average fusaric acid content of 37 mg/kg, compared to 112 mg/kg in kernels only containing fusaric acid. Fusaric acid inhibits plant growth, and is suggested to be antifungal and antinematodic^[201,202]. The high concentration in samples without fumonisins, indicate that non-toxigenic *Fusaria* are present in maize, and supports the possibility of using these fungi as biocontrol agents against fumonisin producing species, as is the case for aflatoxins and *A. flavus*^[203-205].

Equisetin is a known metabolite from *F. equiseti*^[206], which also produce nivalenol^[147]; however, this species was not isolated during the fungal screening. Nivalenol could also originate from other *Fusarium* species like *F. graminearum*^[147], although this species was only seen once in the screening of 240 kernels. The domination of *F. subglutinans* and *F. verticillioides* is most likely the cause for suppression of other *Fusarium* species.

Emodin is produced regularly by several *Eurotium* and *A. wentii*, they are common superficial contaminants of dried grain. The finding of these compounds in the visually uninfected kernels, could be caused by superficial growth, which is easy abraded off the kernels, that hereby appear uninfected. Another important mycotoxin in African maize is aflatoxin^[207], yet no aflatoxin was detected in either of the kernels, which is an outcome of the absence of *A. flavus*/*A. parasiticus* in the fungal screening.

The low amount of fumonisin positive kernels led to the thought of how the fumonisin concentration would be affected if the poor-quality kernels were sorted out from the batch. This effect were estimated by removal of the 4% highly infected kernels (>100 mg/kg). A 71% reduction in fumonisin levels was estimated, demonstrating that this simple approach is a powerful tool for mycotoxin reduction. The strategy of removing infected kernels was successfully implemented in an intervention study in the same rural area of Transkei^[208]. Training of the farmers led to a better result than estimated, giving an 84% reduction with less than 4% weight loss^[198,208,209].

3.4 Fumonisin production by other fungal genera

For almost 30 years *Fusarium* was known as the only fumonisin producer, and then came the discovery of a production of FB₂, FB₄ and the newly discovered FB₆ in *A. niger*^[34-36]. This small group of fumonisin producing fungi was further expanded during a routine screening of cyclosporine producing fungi with a multi-detection method^[199], that led to the discovery that a *Tolypocladium* (figure 9) strain also produced fumonisins. *Tolypocladium* are used in the medical industry for the production of cyclosporines. The cyclosporines are used worldwide as a immunosuppressive drug in organ transplant recipients^[210].



Figure 9: *Tolypocladium inflatum* on YES agar after 7 days growth at 25°C

Due to the important use of *Tolypocladium* species, further investigation of this new exiting discovery was performed. Screening of 12 strains from three different species showed that all 12 produced FB₂, but only ten produced FB₄, table 17. The additional FB's (FB₁, FB₃, FB₆) were not detected in the 12 strains.

Table 17: Production of FB₂ and FB₄ by *Tolypocladium cylindrosporum*, *T. geodes* and *T. inflatum*

Species	FB ₂ positive	FB ₄ positive
<i>T. cylindrosporum</i>	3/3	3/3
<i>T. geodes</i>	3/3	1/3
<i>T. inflatum</i>	5/5	5/5

The fumonisin profile of *Tolypocladium* is similar to that of *A. niger*, but significantly different from that of *Fusarium*. This appear to be in agreement with the proposed fumonisin biosynthesis of Butchko *et al.*^[215], where FB₂ and FB₄ are products from the same biosynthetic pathway. FB₁ and FB₃ are produced in a different pathway, which indicate that *Tolypocladium*, just as *A. niger*, are at least missing the *FUM2* gene in the fumonisin gene cluster^[215].

To obtain new details on the fumonisin producing *Tolypocladium* species, the effect of media composition and temperature was investigated. Three strains were chosen and inoculated on ten different laboratory media along with an *A. niger* strain as a control organism. The ten media were CYA^[166], CYAS^[166], dichloran Rose bengal yeast extract sucrose agar (DRYES)^[211], potato carrot agar (PCA)^[212], malt extract agar (MEA according to Blakeslee)^[147], oatmeal agar (OAT)^[166], PDA (Difco), V8-juice agar with antibiotics (V8)^[212], YES^[166], and dichloran 18% glycerol agar (DG18)^[213]. On each medium at least one of the *Tolypocladium* strains were able to produce fumonisins, figure

10. Two media, PDA and YES, stood out as the best for all three strains. Levels of FB₄ to FB₂ were in the range of 0-95%; however, mostly between 10-30%. The maximal total fumonisin production (FB₂ + FB₄) of the three strains was 0.072 µg/cm² for IBT 41581, 30 µg/cm² for IBT 41582, and 1.3 µg/cm² for IBT 41583. These values are similar to the earlier study (Chapter 3.1) with *Fusarium* and *A. niger* and their detected amounts of 0.006-22 µg/cm² and 0.27-21 µg/cm², respectively.

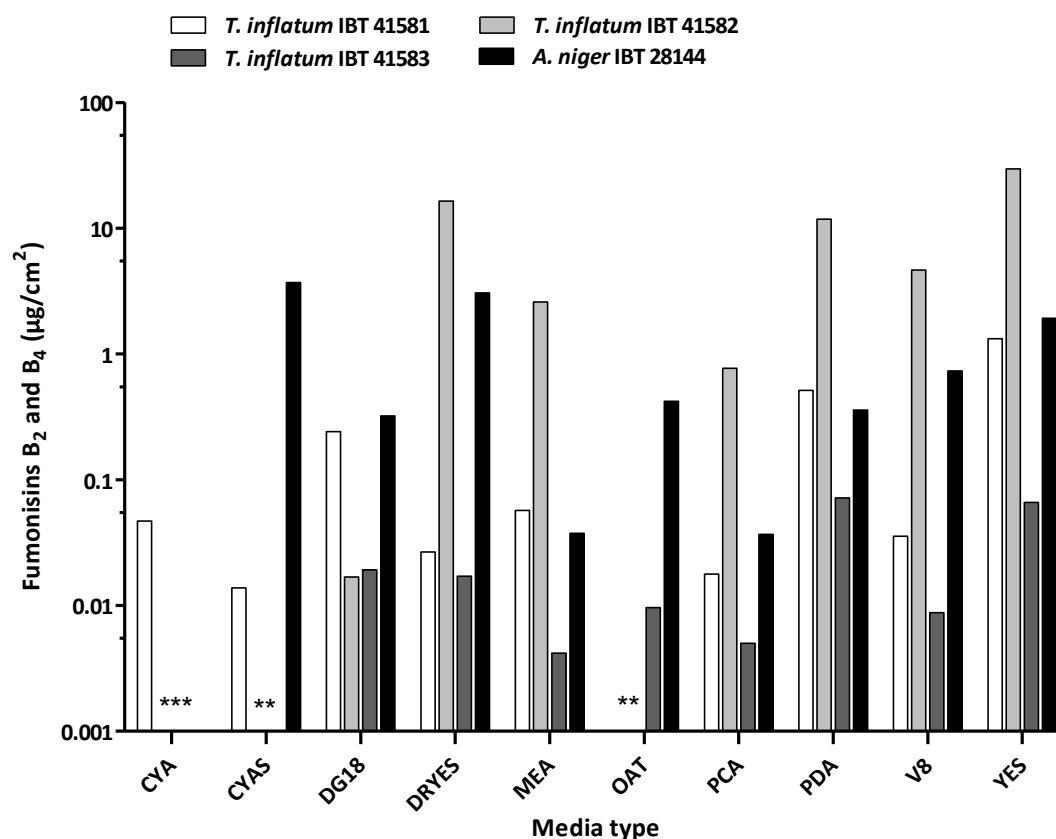


Figure 10: Total production of fumonisin (FB₂ and FB₄) by three *Tolypocladium inflatum* strains (IBT 41581, IBT 41582, IBT 41583) and one *Aspergillus niger* (IBT 28144) on different media at 25°C. *Not detected

To determine further criteria for fumonisin production by *Tolypocladium*, the effect of temperature were studied. Five different temperatures in the range of 15-37 °C were used. Fumonisin production occurred from 15-30 °C, figure 11; samples at 37 °C were not analyzed due to the inability to grow at this temperature. The temperature effect appeared to be strain dependent, since two strains had a maximal production of fumonisins at 30 °C and the last at 25 °C. Even though *Tolypocladium* is not a danger in food, there are possibilities for a fumonisin contamination from this fungus. Not only is it used in the pharmaceutical industry as a producer of cyclosporin A^[210], but it has also been suggested as a possible fungal biological control agent, because of the entomopathogenicity of cyclosporines and efraeptins^[214]. The in house certified

cyclosporin A standard was tested, but did not contain any detectable fumonisins. With this new discovery, the amount of reported fumonisin producing organisms is expanded to a total of 31 species, Table 2.

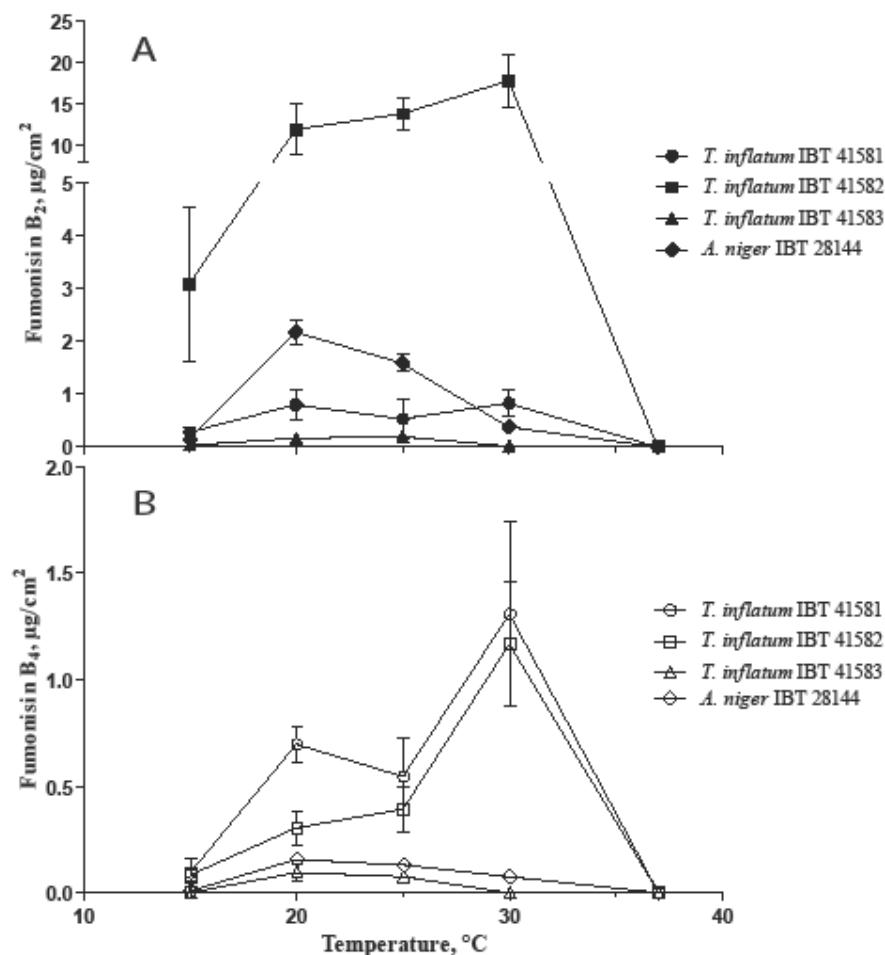


Figure 11: Production of fumonisins B₂ (A) and B₄ (B) by three strains of *Tolypocladium inflatum* (IBT 41581, IBT 41582, IBT 41583) incubated for 7 days at 15, 20, 25, 30 and 37 °C on YES. The standard deviation of triplicates is shown by the error bars. Fumonisin concentration was 0 µg/cm² at 37 °C due to lack of growth.

3.5 Genotoxicity studies of fumonisins

Some work during the past three years has been put into the investigation of the genotoxicity of fumonisins, as seen in chapter 2. The overall idea of this project was to purify FB₂, FB₄, and FB₆ from *A. niger* and compare the genotoxicity of these compounds to FB₁ and FB₂ from *Fusarium*. Since the purification process was difficult to carry out, only a limited amount of preliminary work was performed. The assay of choice, was the comet assay, also called single cell gel electrophoresis. This assay is widely regarded as a quick and reliable method for analysis of DNA

damage in individual cells^[216]. Optimization of the assay, as well as preliminary testing of fumonisins was carried out. Fumonisins were tested against two cell lines, HepG2 (liver) and Caco-2 (colon). Fumonisin B₁ and FB₂ from *Fusarium* were only tested in single replicates at concentrations up to 588 µg/ml, with no signs of either cytotoxicity or genotoxicity; however, these results and replicates are too few to make any firm conclusions on. It is noteworthy that available information of the genotoxic properties of FB₁ is limited and controversial due to conflicting results^[31,217]. This sub-project stalled, since the purification process of fumonisins from *A. niger* was problematic, often resulting in methylated fumonisins, but the purification is still ongoing. Next step in this, is the making of a mutant with an over-expression of the *FUM* genes.

3.6 Overall discussion

Aspergillus niger produced fumonisins in every single tested commodity, raisin, grapes, and peanuts. However the reported fumonisin levels in the food and feed from the past three years are all within the range of 1-25 µg/kg, with one exception of 400µg/kg in a grape must sample^[36,113,114,137,218]. The levels in raisins reported by Varga *et al.* appear to be doubtful due to the high levels and the reported FB₁ and FB₃ production^[136], it is furthermore disputed^[219,220] and is therefore excluded from this comparison. The levels of fumonisins are significantly lower than those produced in worst-case scenarios, indicating that fumonisins from *A. niger* are not a significant problem, since the highest concentration found is 10-160 times (omitting the grape must sample) below the regulatory limit for fumonisins in maize and maize products set by EU^[37]. This is most likely due to the enormous effort put into reducing the already known problem, with ochratoxin A in similar commodities. The visual screening in the field as well as mechanical sorting applications like color sorters^[24], has significantly reduced the mycotoxin contamination. Multiple washing steps are also implemented in the production of certain food, e.g. raisins^[184]; washing crops could also be the cause for the low contamination since this procedure have previously shown to reduce the mycotoxin concentration^[198,221]. Furthermore, implementation of “Good Agricultural Practice” and “Code of Practice” from FAO/WHO^[222] and similar organizations is also likely a liable reason for the significant impact within prevention and reduction of mycotoxin contamination in food and feeds.

Due to the status as an industrial workhorse, production of fumonisins by *A. niger* in an industrial setting, is essential to investigate. Three industrial strains (NRRL 3= IBT 23539; NRRL 567=IBT 26387, NRRL 2001=IBT 26392) used in this study have previously been used for production of organic acids and extracellular enzymes^[223-226]. All three strains produced fumonisins on solid media under all conditions, except high temperature (>37 °C). A recent report mimicked industrial citric acid production conditions and tested *A. niger* production of fumonisins and ochratoxin A. In most cases fumonisins as well as ochratoxin were produced^[146]. In this context, it should be mentioned that industrial production companies, presumably have knocked the fumonisin gene cluster out of their *A. niger* production strains. If this is not the case, fumonisin production could likely occur during industrial conditions, and monitoring industrial end products and/or waste being used as feed for domestic animals^[227,228] is essential. Yet, it should be mentioned that batches of all industrial products are tested for cytotoxicity, carcinogenicity, and similar diverse effects before product approval. These tests would have indicated if ochratoxins and/or fumonisins are produced, but no reports of this have been published^[154].

Preliminary results from the genotoxicity study reveal that FB₁ and FB₂ from *Fusarium* is not genotoxic, in concordance with other studies^[229,230]. However fumonisins from *A. niger* is not tested yet, and this could change the picture completely due to e.g. stereochemistry. Because several genes are not present in the fumonisin gene cluster compared to that of *Fusarium*^[55,56], a different biosynthesis could occur and hereby cause differences in the stereochemistry. An example of the vital role of stereochemistry is thalidomide, where small changes in the stereochemistry made a harmless compound active^[231]. Currently, there are strong indications that the stereochemistry of FB₂ originating from *Fusarium* and FB₂ from *A. niger* is similar^[35]. Additionally, the toxicity of FB₄ and FB₆ has never been tested, even though they are produced to a lesser extent than FB₂, they could still be more potent. Toxicity combinations of mycotoxins are also imperative to investigate, since mycotoxins most likely are present in combinations, rather than alone. With the discovery of a fumonisin production in *A. niger* the combination of ochratoxin and fumonisins is even more relevant, especially because more than 60 % of ochratoxin A producing *A. niger* also produces fumonisins^[146].

4 Conclusion

The frequency of fumonisin producers within the *A. niger* species is approx. 75%, and fumonisins are produced in grapes, raisins, and peanuts at high levels. Although, the frequency and level of fumonisins in retail products are significantly lower. Grapes appear to be the most central commodity for the fumonisin production, affecting the derived products raisins and wine where fumonisins produced by *A. niger* could be detected in 48 % and 23 % of the investigated retail products.

The overall objective of the project is to determine if fumonisins from *A. niger* is a health risk. On the basis of this thesis, there is no evidence that suggest that fumonisins from *A. niger* constitutes a health risk. Especially compared to fumonisins produced by *Fusarium* in maize, since they are found at considerably higher levels.

There is still a small uncertainty, since the toxicity of FB₂, FB₄ and FB₆ from *A. niger* is yet to be evaluated. If the toxicity is similar to fumonisins from *Fusarium*, the significance of this fungi/mycotoxin combination is low, due to the low almost insignificant amounts of fumonisins from *A. niger* in retail products. But at the present moment fumonisins from *A. niger* should not be considered a health risk.

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6 Original manuscripts (1-9)

- Paper 1:** J.M. Mogensen, K.F. Nielsen, R.A. Samson, J.C. Frisvad and U. Thrane. Effect of temperature and water activity on the production of fumonisins by *Aspergillus niger* and different *Fusarium* spp. BMC Microbiology. 2009. 9:281
- Paper 2:** J.M. Mogensen, J.C. Frisvad, U. Thrane, K.F. Nielsen. Production of fumonisin B₂ and B₄ by *Aspergillus niger* on grapes and raisins. Journal of Agricultural and Food Chemistry. 2010, 58:954-958
- Paper 3:** J.M. Mogensen, T.O. Larsen, K.F. Nielsen. Widespread occurrence of the mycotoxin Fumonisin B₂ in wine. Journal of Agricultural and Food Chemistry. 2010, 58:4583-4587
- Paper 4:** P. B. Knudsen, J. M. Mogensen, T. O. Larsen and K. F. Nielsen. Occurrence of fumonisins B₂ and B₄ in retail raisins. Journal of Agricultural and Food Chemistry. 2011, 59:772-776
- Paper 5:** J.M. Mogensen. J. Varga, U. Thrane and J.C. Frisvad. *Aspergillus acidus* from Puerh tea and black tea does not produce ochratoxin A and fumonisin B₂. International Journal of Food Microbiology. 2009. 132:141-144.
- Paper 6:** J. M. Mogensen, E. O. Søndergaard and K. F. Nielsen. Studies on fumonisins in peanuts and peanut butter. Journal of Agricultural and Food Chemistry. Submitted.
- Paper 7:** J. M. Mogensen, S. M. Sørensen, M. Sulyok, L. van der Westhuizen, G. Shephard, J.C. Frisvad, U. Thrane, R. Krska, K. F. Nielsen. Single kernel analysis of fumonisins and other fungal metabolites in maize from South African subsistence farmers. Food Additives & Contaminants. 2011. 28:1724-1734.
- Paper 8:** J. M. Mogensen, K. A. Møller, P. von Freiesleben, R. Labuda, E. Varga, M. Sulyok, A. Kubátová, U. Thrane, B. Andersen, K. F. Nielsen. Production of Fumonisin B₂ and B₄ in *Tolypocladium* species. Journal of Industrial Microbiology and Biotechnology. 2011, 38:1329-1335.
- Paper 9:** K.F. Nielsen, J.M. Mogensen, M. Johansen, T.O. Larsen and J.C. Frisvad. Review of secondary metabolites and mycotoxins from the *Aspergillus niger* group. Analytical & Bioanalytical Chemistry. 2009. 395:1225-1242.

Paper 1

” Effect of temperature and water activity on the production of fumonisins by *Aspergillus niger* and different *Fusarium* species”

J.M. Mogensen, K.F. Nielsen, R.A. Samson, J.C. Frisvad and U. Thrane

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Research article

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Effect of temperature and water activity on the production of fumonisins by *Aspergillus niger* and different *Fusarium* species

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Abstract

Background: Fumonisins are economically important mycotoxins which until recently were considered to originate from only a few *Fusarium* species. However recently a putative fumonisin gene cluster was discovered in two different *Aspergillus niger* strains followed by detection of an actual fumonisin B₂ (FB₂) production in four strains of this biotechnologically important workhorse.

Results: In the present study, a screening of 5 *A. niger* strains and 25 assumed fumonisin producing *Fusarium* strains from 6 species, showed that all 5 *A. niger* strains produced FB₂ and 23 of 25 *Fusarium* produced fumonisin B₁ and other isoforms (fumonisin B₂ and B₃). Five *A. niger* and five *Fusarium* spp. were incubated at six different temperatures from 15-42°C on Czapek Yeast Agar +5% salt or Potato Dextrose Agar. *A. niger* had the highest production of FB₂ at 25-30°C whereas *Fusarium* spp. had the maximal production of FB₁ and FB₂ at 20-25°C. Addition of 2.5-5% NaCl, or 10-20% sucrose increased the FB₂ production of *A. niger*, whereas addition of glycerol reduced FB₂ production. All three water activity lowering solutes reduced the fumonisin production of the *Fusarium* species.

Conclusion: The present study shows that the regulation of fumonisin production is very different in *A. niger* and *Fusarium*, and that food and feeds preserved by addition of sugar or salts may be good substrates for fumonisin B₂ production by *A. niger*.

Background

The fumonisins were discovered in 1988 and are divided in four series A, B, C, P [1-3] with the B₁ (FB₁), B₂ (FB₂) and B₃ (FB₃) as the most abundant naturally occurring homologues [4,5]. They were first isolated from *Fusarium verticillioides* (= *F. moniliforme* pro parte [6]) strain MRC 826 by Gelderblom *et al.* [7]. FB₁ is mainly produced by *F. verticillioides* and *F. proliferatum* [8]. However, production of type

B fumonisins by other *Fusarium* spp. has also been reported, e.g. from *F. dlamini*, *F. napiforme*, *F. nygamai* and *F. oxysporum* [8-10]. Fumonisins are important mycotoxins because they are suspected to cause human and animal toxicoses by the consumption of contaminated corn-based food and feeds [11]. Fumonisins have been shown to induce outbreaks of equine leukoencephalomalacia in horses and pulmonary edema and hydrothorax in pigs

[5,12]. The fumonisins are structurally similar to sphingolipids and have shown to inhibit the sphingolipid biosynthesis via the ceramide synthase pathway [13,14]. To avoid possible health risks, the U.S. Food and Drug Administration recommends that corn products should not be used for human consumption when contaminated with more than 2-4 mg/kg total fumonisins (depending on the product) [15], whereas EEC has a regulatory limit of 0.2-2 mg/kg (depending on the product) [16].

Fumonisins produced by *Fusarium* species have been isolated from corn [1] and corn based products [11] such as tortillas [17] and beer [18], as well as other commodities like rice [19], black tea leaves [20], asparagus [21] and pine nuts [22].

Factors that affect the production of fumonisins in *Fusarium* have been well studied, and include solid substrates [23], liquid substrates [24], temperature [25-27], water activity (a_w) [27,28], pH [29], addition of nitrogen repressor [30], aeration of the substrate [29] and addition of fumonisin precursors [31], but often corn kernels have been used as substrate since corn is the primary crop infected with *F. verticillioides*.

Recently putative homologues to the *F. verticillioides* fumonisin gene cluster were found in two different *Aspergillus niger* genomes [32,33], and it was subsequently shown that three full genome sequenced strains and the ex type strain of *A. niger* actually can produce FB₂ in comparable amounts to *Fusarium* strains [34] when grown on agar substrates with high amounts of sugar, glycerol or NaCl.

This was followed by the discovery of additional FB₄ production (~20% the amount of FB₂) by *A. niger* [35] in agar cultures and naturally *A. niger* contaminated Thai coffee beans [35].

The objectives of the present work were to i) screen *A. niger* and *Fusarium* strains, for production of FB₁, FB₂ and FB₃ on three different agar substrates, ii) investigate the effect of incubation temperature on the production of fumonisins and iii) study the effect of the solutes glycerol, NaCl and sucrose on the production of fumonisins. The current work is performed on agar media instead of natural substrates in order to more easily assess the water activity.

Results

Optimization of extraction

The efficiency of five different extraction solvents to extract FB₂ from *A. niger* (NRRL 567) varied significantly, with methanol:water (3:1) being most efficient, followed by acetonitrile:water (3:1) with a 20% lower efficiency

and methanol:dichloromethane:ethyl acetate (1:2:3) (30% lower efficiency). The use of water (25°C) and hot water (100°C) was not suitable for extraction of FB₂ from *A. niger* NRRL 567 with a relative efficiency of <1%, when compared to methanol:water. The most effective solvents concerning *F. verticillioides* IBT 9400 were acetonitrile:water (3:1) as the best followed by methanol:water (3:1) (98%) and water at room temperature with an efficiency of 93%. The use of hot water as extraction solvent was less efficient (76%), compared to methanol:water. The acidic methanol:dichloromethane:ethyl acetate (1:2:3), had the lowest extraction efficiency with 26%. The difference between each of the two replicates for all incidences was in the interval 1-40% with an average of 10%.

Validation of methanol-water extraction from *A. niger*

The recovery of FB₂ from two spiked non-FB₂ producing strains showed a recovery of 75% ± 10% (IBT 20381) and 85% ± 10% (IBT 19345). The calibration curves from standards and spiked samples, used to calculate recovery from all had R² better than 0.995. The relative standard deviation (RSD) of the extracted amounts of FB₂ from the 8 isolates (n = 5) varied within 4-50%, with an average RSD of 20%. LOD were found to be 0.1 µg/cm² fungal culture.

Screening of strains for fumonisin production

The results from the screening experiment are shown in table 1. The *A. niger* strains were able to produce FB₂ on all three substrates, with the highest production on RC and CYAS. None of the *A. niger* strains produced detectable amounts of neither FB₁ nor FB₃. LC-MS/MS analyses have shown that FB₂ is produced along with FB₄, although the amount of FB₄ normally lies in the range 5-20% of the FB₂ amounts [35].

Of the *A. niger* strains, NRRL 567 had the highest production of FB₂ on RC and CYAS. But on PDA the three strains, NRRL 567, NRRL 2001 and IBT 24631, produced very similar amounts of fumonisins, although lower than RC and CYAS. The FB₂ production on RC and CYAS of *A. niger* NRRL 2001, IBT 24631 and 24634 differed only slightly, whereas FB₂ production by the other two, NRRL 567 and NRRL 3 were clearly favored by growth on RC. All *Fusarium* spp. with the exception of the two *F. dlamini* strains IBT 2937 and IBT 2938 produced fumonisins under these conditions. Six *Fusarium* strains, *F. napiforme* IBT 2932, *F. proliferatum* IBT 9109 and IBT 9337, *F. verticillioides* IBT 9496 and *F. nygamai* IBT 9395 produced amounts close to the detection limit of FB₁ on RC; in addition IBT 2932 also produced fumonisins on PDA. *F. nygamai* IBT 2934, IBT 8554 and IBT 8557 showed a higher production of FB₂ than FB₁ on PDA (data not shown), and *F. proliferatum* IBT 9397 had the highest concentration of total fumonisin

Table 1: Fumonisin production by *Aspergillus niger* and *Fusarium* spp. on CYAS, PDA and RC after 7 days growth at 25°C.

Fungi	Isolate	CYAS g/ml	RC g/ml	PDA g/ml
<i>A. niger</i>	NRRL 3	2.9 ± 0.4	7.9 ± 0.7	0.86 ± 0.02
	NRRL 567	25 ± 0.9	36 ± 2	1.9 ± 0.5
	NRRL 2001	7.6 ± 0.7	6.1 ± 0.9	3.1 ± 0.3
	IBT 24631	5.2 ± 0.2	6.7 ± 0.7	1.3 ± 0.6
	IBT 24634	6.4 ± 0.1	5.3 ± 0.4	0.46 ± 0.02
<i>F. proliferatum</i>	IBT 8904	n.d.	9.9 ± 3	21 ± 0.5
	IBT 9109	n.d.	0.028 ± 0.007	n.d.
	IBT 9337	n.d.	0.021 ± 0.01	n.d.
	IBT 9393	n.d.	0.03 ± 0.001	2.0 ± 0.04
	IBT 9397	n.d.	46 ± 3	33 ± 0.5
	IBT 41107	n.d.	5.4 ± 1	6.7 ± 0.6
<i>F. verticillioides</i>	IBT 9400	n.d.	0.035 ± 0.005	35 ± 0.7
	IBT 9492	n.d.	0.028 ± 0.004	4.9 ± 0.2
	IBT 9496	n.d.	0.033 ± 0.0003	n.d.
	IBT 9502	n.d.	2.2 ± 0.4	18 ± 1
	IBT 9505	n.d.	0.078 ± 0.09	9.5 ± 3
	IBT 41110	n.d.	0.12 ± 0.07	4.5 ± 0.7
<i>F. dlamini</i>	IBT 2937	n.d.	n.d.	n.d.
	IBT 2938	n.d.	n.d.	n.d.
<i>F. napiforme</i>	IBT 2931	n.d.	0.24 ± 0.06	6.2 ± 0.9
	IBT 2932	n.d.	0.13 ± 0.1	0.081 ± 0.02
<i>F. nygamai</i>	IBT 2933	n.d.	0.033 ± 0.006	n.d.
	IBT 2934	n.d.	22 ± 2	5.6 ± 0.6
	IBT 8290	n.d.	0.041 ± 0.006	0.047 ± 0.001
	IBT 8554	n.d.	0.033 ± 0.01	0.039 ± 0.003
	IBT 8557	n.d.	0.01 ± 0.01	0.14 ± 0.002
	IBT 8566	n.d.	3.1 ± 1	6.2 ± 0.5

Table 1: Fumonisin production by *Aspergillus niger* and *Fusarium* spp. on CYAS, PDA and RC after 7 days growth at 25°C. (Continued)

	IBT 9394	n.d.	0.30 ± 0.08	n.d.
	IBT 9395	n.d.	0.03 ± 0.003	16 ± 0.6
<i>F. oxysporum</i>	IBT 9514	n.d.	2.8 ± 0.4	37 ± 0.8

The concentration of FB₂ was detected in the methanol:water (3:1) extract. The values are means of the replicates plus/minus the standard deviation. The replicates are made in biological duplicates on two separate plates.

n.d. not detected.

Standard deviation calculated on two measurements

measured. Only seven strains had a total production of fumonisin above 1 µg/ml extract on RC compared to 14 strains on PDA.

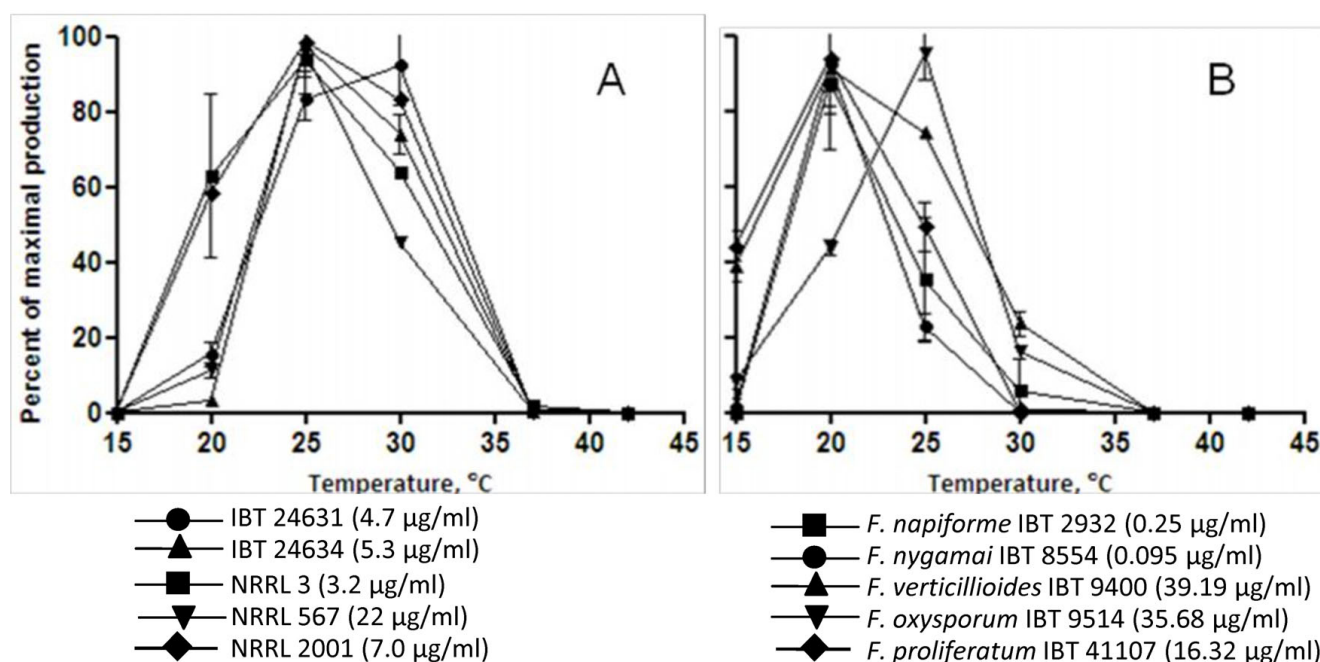
Seven *Fusarium* strains did not have any measurable production of FB₁, FB₂ nor FB₃ on PDA compared to two non-producers on RC. On the other hand PDA supported production of higher amounts of fumonisins, strains which had a barely detectable amount of FB₁ on RC, also showed production of either FB₂ or both FB₂ and FB₃ on PDA.

Strains for the next experiments were selected on the basis of the above mentioned experiments. Besides the five *A.*

niger strains five *Fusarium* strains were selected, with both good and poor producers at 25°C, two strains with a high FB production, *F. verticillioides* IBT 9400 and *F. oxysporum* IBT 9514, one with an average FB production *F. proliferatum* IBT 41107, one with a low FB₁ production *F. napiforme* IBT 2932 and a strain with a higher production of FB₂ than FB₁ *F. nygamai* IBT 8554 was selected.

The effect of temperature on growth and production of fumonisins by *A. niger* and *Fusarium* spp

Only one of the *A. niger* strains was able to grow at 15°C, although very slowly (extensive data shown in [Additional file 1]). The growth increased at higher temperature

**Figure 1**

A: FB₂ production by *Aspergillus niger* after 7 days of growth at different temperatures (15-42°C) on CYAS. The concentration of FB₂ was detected in the methanol:water (3:1) extract. The values are means of biological duplicates on two different plates; highest value (µg/ml) is in parenthesis. **B: Total production of fumonisin B₁, B₂ and B₃ produced by *Fusarium* spp. after 7 days growth at different temperatures (15-42°C) on PDA.** The concentration of fumonisin was detected in the methanol:water (3:1) extract. The values are means of biological duplicates on two different plates; highest value (µg/ml) is in parenthesis.

and peaked at 30-37°C, followed by a slight reduction of the growth at 42°C. All *Fusarium* strains were able to grow at 15°C followed by an increased growth at higher temperatures and peaked at 25-30°C, above this temperature the growth decreased and no growth was observed at 42°C. The effect of temperature on the production of FB₂ by the five *A. niger* strains is shown in figure 1A. None of the isolates produced detectable amounts of FB₂ at 42°C, even though all strains grew well. The only *A. niger* strain, NRRL 2001, that was able to grow at 15°C, did not have any detectable production of FB₂.

At 37°C, the optimal temperature for growth by *A. niger*, the FB₂ production was very limited. NRRL 567 had the highest production of the five strains at 25°C and 30°C; however at 20°C NRRL 2001 showed the highest production. This strain showed also the highest growth rate at this temperature. Four strains, NRRL 567, NRRL 3, NRRL 2001 and IBT 24634 had the highest FB₂ production at 25°C, followed by 30°C, 20°C, and 37°C (Figure 1A). One strain, IBT 24631, differed from the other four, since it had a maximal production at 30°C, followed by 25°C, 20°C and 37°C.

The concentration of FB₁, FB₂ and FB₃ produced by *Fusarium* spp. is shown in figure 1B, and four of the *Fusarium* spp. had maximal production at 20°C: *F. verticillioides* IBT 9400, *F. proliferatum* IBT 41107, *F. napiforme* IBT 2932 and *F. nygamai* IBT 8554. *F. oxysporum* IBT 9514 had a maximal production of fumonisin at 25°C. Only three of the *Fusarium* strains, *F. verticillioides* IBT 9400, *F. oxysporum* IBT 9514 and *F. proliferatum* IBT 41107 had a measurable production of fumonisin at 15°C. *F. napiforme* IBT 2932 was only able to produce detectable amounts of fumonisins in the temperature range 20-25°C. At 20°C this strain had detectable concentrations of FB₁, FB₂ and FB₃, but at 25°C only FB₁ and FB₂ were detected (data not shown). Even though there was growth of all five *Fusarium* strains at 37°C there were only in one case detectable production of fumonisins, this was produced by *F. verticillioides*.

The effect of glycerol, NaCl and sucrose on the growth and production of fumonisins by *A. niger* and *Fusarium* spp

All strains of *A. niger* and *Fusarium* spp. were able to grow at all glycerol concentration (0-255 g/l) [see Additional file 2]. The growth of *A. niger* was only slightly reduced at a_w 0.99; below this there was a continuous decrease in the growth. For the *Fusarium* spp. the growth was reduced at a_w 0.99 and below, which was the same as the *A. niger*. All *A. niger* strains were able to produce FB₂ at all glycerol concentrations tested (Figure 2). The effect of glycerol on the quantitative FB₂ production for *A. niger* was very strain dependent, but in general the average FB₂ production was reduced 8.2% per 0.01 a_w unit ($R^2 = 0.97$). Three of the five *Fusarium* strains were able to produce fumonisin at all

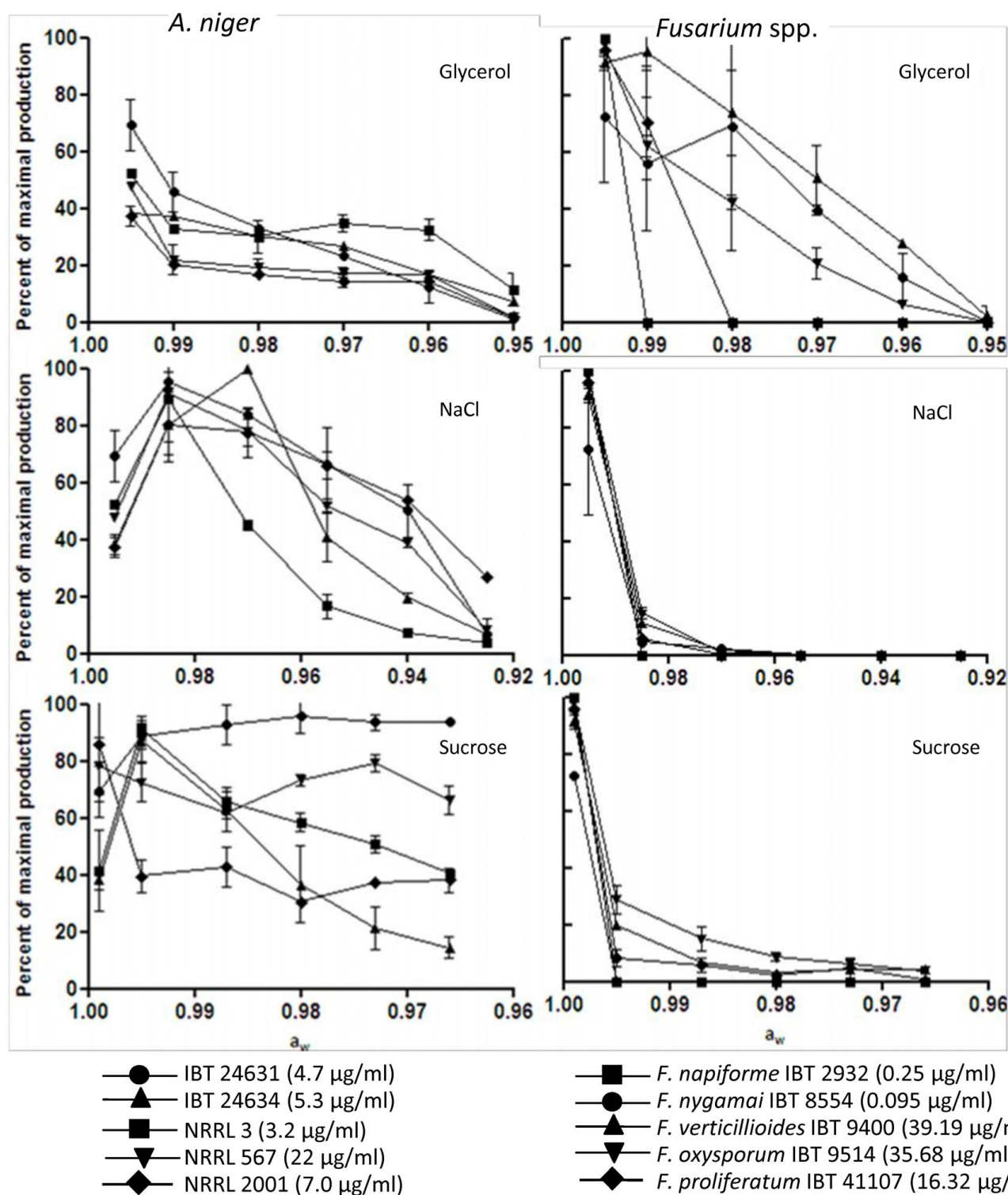
glycerol concentrations: *F. nygamai* IBT 8554, *F. oxysporum* IBT 9514 and *F. verticillioides* IBT 9400 (Figure 2). For two of these, IBT 8554 and IBT 9400, fumonisin production was increased up to 20% when glycerol was added. The other two strains *F. napiforme* IBT 2932 and *F. nygamai* IBT 8554 did not have a measurable production of fumonisins at a_w 0.99 and 0.98. The average total fumonisin production was reduced 18% per 0.01 a_w unit (linear regression $R^2 = 0.91$).

For the five *A. niger* growth in the presence of NaCl was partially inhibited below a_w 0.98 and all strains were able to grow at all a_w tested. The *Fusarium* species had the highest growth rates at the highest a_w and the growth declined until a_w 0.93 where they were unable to grow.

The FB₂ production of all *A. niger* strains was stimulated when NaCl was added with FB₂ being detected at all NaCl concentrations (Figure 2). The production peaked at a_w 0.985-0.97 depending on the strain. A reduction of the FB₂ production started at a_w 0.97-0.94 and a comparison by a regression analysis to the cultures with no NaCl added the average production of FB₂ decreased 13% per 0.01 a_w unit ($R^2 = 0.98$). The addition of NaCl clearly reduces the production of fumonisins by *Fusarium* (Figure 2), and was not detected in any culture below a_w 0.955. One strain, *F. napiforme* IBT 2932, did not even have a measurable production of fumonisin when NaCl was added. But no correlations in the average decrease of fumonisin production was observed.

The *A. niger* and *Fusarium* spp. strains grew better on sucrose, compared to NaCl and glycerol. The addition of sucrose increased the production of FB₂ for all the *A. niger* strains (Figure 2). The increase in the FB₂ production was followed by either a reduction or stagnation in the production level at higher sucrose concentrations. The average decrease in the total fumonisin production were 8.3% per 0.01 a_w unit ($R^2 = 0.87$)

Only three of the five *Fusarium* strains were able to produce FB₁, FB₂ and FB₃ when cultured on different sucrose concentrations (Figure 2). Because a commercial potato extract without sucrose was not obtainable, the base potato extract was prepared on home-made boiled potatoes. The three *Fusarium* strains with a detectable fumonisin production were *F. oxysporum* IBT 9514, *F. proliferatum* IBT 41107 and *F. verticillioides* IBT 9400. *Fusarium oxysporum* IBT 9514 had a maximal production at a_w 0.995, while *F. proliferatum* IBT 41107 peaked at a_w 0.999 and *F. verticillioides* IBT 9400 peaked at a_w 0.999. A reduction of the fumonisin production was observed at higher sucrose concentrations. There were no correlations in the average decrease of fumonisin production

**Figure 2**

Effect of glycerol, NaCl and sucrose on the FB_2 production of *Aspergillus niger* and the total production of FB_1 , FB_2 and FB_3 by *F. verticillioides*, *F. proliferatum*, *F. napiforme*, *F. nygamai* and *F. oxysporum*. Strains have been incubated at 25°C for 7 days on CYA or PDA. The concentration of fumonisins were detected in the methanol:water (3:1) extract. The values are means of biological duplicates on two different plates, highest value is in parenthesis.

Discussion

We found that *A. niger* was able to produce FB₂, in agreement with Frisvad *et al.* [34] who showed that *A. niger* produced FB₂ on the agar substrates RC and CYAS. On the other hand they did not measure any production of FB₂ on PDA, whereas all five *A. niger* strains in the present study had a detectable, albeit low, production of FB₂ on this substrate. Besides *F. verticillioides*, also *F. napiforme*, *F. nygamai*, *F. proliferatum* and *F. oxysporum* were found to produce fumonisins on laboratory agar substrates. The ability to produce fumonisins by these species correlates with findings of Nelson *et al.* [8] and Kpodo *et al.* [9]. However Nelson *et al.* [8] described the production of FB₁ by *F. dlamini*, but this was not supported in our study. The findings of strains, capable of producing more FB₂ than FB₁ was also described by Musser & Plattner [36] as well as Leslie *et al.* [37]. Apart from the four *A. niger* strains shown to produce FB₂ by Frisvad *et al.* [34], additional 12 strains did produce this mycotoxin. Among the 18 *A. niger* strains investigated until now, only two have been unable to produce fumonisins in detectable amounts on the media investigated.

Astoreca *et al.* [38] found the optimal temperature for growth of *A. niger* to be 30°C, the highest investigated temperature in their study. In correlation to this study Palacios-Cabrera *et al.* [39] also found that *A. niger* grew optimally at temperatures of 30°C, which was also the optimal temperature for linear growth in our study. Marin *et al.* [40] found that the growth of both *F. verticillioides* and *F. proliferatum* was best at 25 to 30°C, which is in agreement of our results. According to Marin *et al.* [40] *F. verticillioides* is more tolerant to temperature above 30°C than other *Fusarium* spp., however this was not obvious in our study.

We found that the optimal production of FB₂ by *A. niger* was at 25°C and in one case 30°C. Since there has been only one report of the fumonisin production by *A. niger* we have compared our results to the production of ochratoxin A. Our results correlates with an investigation of Esteban *et al.* [41], who showed that the optimal temperature for production of ochratoxin A by *A. niger* was at 20-25°C. In contrast to this, other authors found that the optimal ochratoxin A production in a synthetic grape juice medium was significantly better at 15°C compared to both 25 and 35°C [42]. Findings in our study showed that at 20°C a significant decrease in the FB₂ production occurred compared to 30°C. Earlier studies show that the optimal temperature for production of fumonisins by *F. proliferatum* is at 15-20°C where *F. verticillioides* prefers the higher temperatures of 30°C [27]. This partly correlates with the results from our study, where both isolates showed the highest production at 20°C, but also produced fumonisins at 30°C. However fumonisin produc-

tion by *F. verticillioides* was less inhibited than *F. proliferatum* by the higher temperature. Dilkin *et al.* [26] and Alberts *et al.* [25] found the optimal temperature for fumonisin production to be 25°C, followed by 20 and then 30°C. These results deviate from our results, because four isolates had the best production at 20°C and one at 25°C. Marin *et al.* [27] described the production of FB₁ by *F. verticillioides* at 37°C, which was also observed in our study. From our results a general pattern in the fumonisin production for both genera was observed, namely the maximal production of fumonisins being 5°C below the optimal growth.

Comparing *A. niger* with the *Fusarium* spp., all *A. niger* strains grew better at all tested a_w values. Leong *et al.* [42] found the optimal a_w for ochratoxin A production by *A. niger* to be a_w 0.95, whereas Esteban *et al.* [43] found it to be in the range of 0.96-0.99, and that it was very strain dependent. These values are lower than those observed in our study for fumonisins where four of the *A. niger* strains had the highest production of FB₂ at a_w 0.99, with one isolate produced most FB₂ at a_w 0.98. Earlier studies have shown that the optimal a_w for fumonisin production by *Fusarium* is in the interval 0.97-0.98 [27,44]. The optimal a_w value from our study was a_w 0.995, which is a bit higher than the above mentioned a_w values. Frisvad *et al.* [34] also found that the addition of 5% NaCl (a_w = 0.97) or 20% sucrose (a_w = 0.99) increased the production of FB₂ by *A. niger*. The present study showed that not all the used strains had the same pattern. Only four of the five strains had an increase in FB₂ production at a_w 0.97 compared to the zero sample when grown on NaCl. The last *A. niger* NRRL 3 had only an increase at a_w 0.985 followed by a decrease at higher NaCl concentrations. The same was observed with sucrose where three strains had an increase in the FB₂ production compared to a sucrose concentration of 3% in standard medium. The last strains had a decrease in the production at the previous mentioned sucrose concentrations. In conclusion it is clear that there is very large strain variability in fumonisin production at different water activities between the *A. niger* strains used in these experiments.

Further studies on the effect of physiological variables on fumonisin production by *A. niger* is needed on large numbers of strains, due to large strain differences, and should further be backed up by studies of commodities where *A. niger* is common. This will lead us to a better understanding of how large a food safety problem fumonisin production by *A. niger* is. Only one report of fumonisins from *A. niger* in food (green coffee beans) has been reported, however the amounts were well below the regulatory limit [35]. Fumonisin contamination in a series of foods, often contaminated with *A. niger*, may show to be an overlooked mycotoxin problem, since surveys have only

mostly targeted corn and rice, based on the belief that it was only *Fusarium* spp. that could produce these important mycotoxins.

Conclusion

The present study shows that the regulation of fumonisin production is very different in *A. niger* compared to *Fusarium*, with the latter preferring high a_w (>0.99) and low temperature (20-25°C) and with *A. niger* preferring lower a_w and higher temperature (25-30°C). Fumonisin produced by *A. niger* may be an overlooked health risk in foods not previously associated with fumonisins, especially because *A. niger* is known as a common food spoilage organism on a wide variety of foods [45].

Methods

Unless otherwise is stated all solvents were HPLC grade, chemicals were analytical grade and water was purified on a Milli-Q system (Millipore, Bedford, MA). Media were prepared in 9 cm Petri dishes, each with 20 ml medium, and strains were inoculated by single point and dishes were incubated in micro perforated plastic bags at 25°C for 7 days in darkness. The colony diameter measured is an average of the smallest and broadest diameter. All samples were as a minimum made in biological duplicates on two individually plates.

Fungal strains and media

All strains (Table 2) were from the IBT culture collection at Center for Microbial Biotechnology, DTU. The *Fusarium* strains were selected from species known to produce fumonisin. Five *A. niger* strains were selected, for the physiologically study, three of these have been used in industry according to collection databases (NRRL 3, NRRL 567 and NRRL 2001), whereas the last two were isolated from black pepper (IBT 24634, IBT 24631). The last 9 strains were only used for validation. All *A. niger* strains were characterized and identified using a polyphasic approach, as in our previous articles [46,47], in order to ensure that they were *A. niger sensu stricto*. Besides this the ITS region and parts of the β -tubulin and calmodulin genes were amplified and sequenced as described previously [48-50]. Results shown in Table 2 (public database numbers of the strains).

The media used for fumonisin production were: potato dextrose agar (PDA) [51], Czapek yeast autolysate agar with 5% NaCl (CYAS) [52] and rice meal corn steep liquor (RC) agar [53].

Fumonisin analysis

The fumonisin were extracted using the method previously described by Frisvad *et al.* [34]. Six plugs (D = 6 mm) were cut out of the colony from the center and in a radius towards the edge of the colony and transferred to a clean

2-ml vial, 800 μ l of methanol:water (3:1) was added, and extracted by ultrasonication for one hour. All extracts were filtered through a 13 mm PTFE 0.45 μ m syringe filter (National Scientific, Rockwood, Tennessee) into a new vial and used directly for analysis.

The LC-MS analysis was performed on a LC/MSD VL single quadrupole (Agilent, Santa Clara, California). The separation of 3 μ l extracts were done at 40°C on a 50 \times 2 mm, i.d. 3 μ m size, Luna C-18 (II) column (Phenomenex, Torrance, California), fitted with a security guard column, and using a water:acetonitrile (both containing 20 mM formic acid) gradient at a flow rate of 0.3 ml/min. The gradient started at 30% acetonitrile, and increased to 60% acetonitrile over 5 minutes. During further 1 minute it was increased to 100% acetonitrile and maintained here for 2 minutes before the gradient in 1 minute was returned to starting conditions and kept there for 5 minutes. The mass spectrometer (MS) was operated in positive electrospray ionization mode and was automatically calibrated on the instrument ESI tuning mix. The MS was used in selected ion monitoring (SIM) mode for measuring $[M+H]^+$: FB₁ (m/z 722), FB₂ and FB₃ (m/z 706). The capillary voltage was held at 3000 V, the fragmentor voltage was at 70 V and the nebulizer pressure was at 2.5 bar. The drying gas flow was 12 l/min with a temperature of 350°C. The detection limit was measured to 0.01 μ g/ml from dilutions of a FB₁ and FB₂ certified standard (Biopure, Tulin, Austria), with concentrations of 50.2 μ g/ml and 51.0 μ g/ml, respectively. The fumonisin concentrations of the extracts were calculated from a standard curve created from dilutions of the FB₁ and FB₂ standard mixture. FB₂ presence was further confirmed in selected extracts by LC-MS/MS [35] and LC-HRMS [34].

Efficiency of extraction solvents

The five different extraction solvents tested on *A. niger* NRRL 567 and *F. verticillioides* IBT 9400 were: i) methanol:water (3:1), ii) acetonitrile:water (3:1), iii) water at room temperature (25°C), iv) water at 100°C and v) methanol:dichloromethane:ethyl acetate (1:2:3) with addition of 1% (v/v) formic acid. The extraction process for [i-iv] was the same as mentioned above. For the fifth extraction solvent there were a few extra steps: After ultrasonication the extract was transferred to a new vial and the organic phase was evaporated *in vacuo*. The residue was redissolved by ultrasonication in 500 μ l methanol for 20 minutes. All extracts were filtered through a PTFE 0.45 μ m syringe filter before analysis.

Validation of methanol:water extraction

FB₂ extraction was validated by spiking 5 plugs of two non-fumonisin producing strains of *Aspergillus niger* (IBT 19345 and IBT 20381) with 100 μ l FB₂ standard containing 5000, 2500, 1000, 500 and 100 ng FB₂. After spiking,

Table 2: Fungal isolates used for fumonisin production.

Fungi	Isolate	Genbank numbers
<i>Aspergillus niger</i>	NRRL 3 (ex unknown) (= ATCC 9069, CBS 120.49, IBT 23539) (Full genome sequenced)	FJ639289
	NRRL 567 (ex unknown) (= ATCC 12846, IBT 26387)	GU195638
	NRRL 2001 (ex unknown) (= ATCC 13794, IBT 26392)	GU195639
	IBT 24631 (ex black pepper)	GU195636
	IBT 24634 (ex black pepper)	GU195637
Only used for validation	IBT 4983 (ex unknown) (= CBS 117.80)	GU195632
	IBT 18741 (ex carpet dust)	FJ639294
	IBT 19345 (ex unknown) (= IFO 6082)	GU195633
	IBT 19558 (ex coffee beans)	GU195634
	IBT 20381 (ex coffee beans)	GU195635
	IBT 26774 (ex unknown)	
	IBT 28086 (ex grape)	
	IBT 28104 (ex black pepper)	GU105640
<i>Fusarium dlamini</i>	IBT 2937 (ex plant debris) (= FRC M-1688)	
	IBT 2938 (ex plant debris) (= FRC M-1638)	
<i>F. napiforme</i>	IBT 2931 (ex soil debris from grassland) (= FRC M-1647)	
	IBT 2932 (ex soil debris from grassland) (= FRC M-1646)	
<i>F. nygamai</i>	IBT 2933 (ex unknown) (= FRC M-2376)	
	IBT 2934 (ex root debris from grassland) (= FRC M-2371)	
	IBT 8290 (ex unknown) (= MRC 4373)	
	IBT 8554 (ex unknown)	
	IBT 8557 (ex unknown)	
	IBT 8566 (ex corn kernel)	
	IBT 9394 (ex unknown) (= MRC 3997)	
	IBT 9395 (ex unknown) (= MRC 3998)	
<i>F. oxysporum</i>	IBT 9514 (ex corn kernel)	
<i>F. proliferatum</i>	IBT 8904 (ex yellow onion)	
	IBT 9109 (ex barley)	
	IBT 9337 (ex corn stalk)	
	IBT 9393 (ex unknown) (= MRC 3218)	
	IBT 9397 (ex unknown) (= MRC 3216)	
	IBT 41107 (ex corn)	
<i>F. verticillioides</i>	IBT 9400 (ex unknown) (= MRC 826)	
	IBT 9492 (ex corn kernel)	
	IBT 9496 (ex corn kernel)	
	IBT 9502 (ex corn kernel)	
	IBT 9505 (ex corn kernel)	
	IBT 41110 (ex corn)	

ATCC: American Type Culture Collection, Manassas, VA, USA

CBS: Centraalbureau voor Schimmelcultures, Utrecht, The Netherlands.

FRC: Fusarium Research Center, Penn State University, University Park, Pennsylvania, USA

IFO: Institute for Fermentation, Osaka, Japan.

MRC: South African Medical Research Council, Tygerberg, South Africa.

NRRL: Northern Regional Research Laboratory, Peoria, Illinois, USA.

Table 3: The concentration of glycerol, NaCl and sucrose and the corresponding measured water activity (a_w)

Glycerol (g/l)	0	51	102	154	204	255	---
a_w	1 ± 0	0.99 ± 0.0015	0.98 ± 0	0.97 ± 0.001	0.96 ± 0.001	0.95 ± 0.001	---
NaCl (g/l)	0	25	50	75	100	125	---
a_w	1 ± 0	0.985 ± 0.0006	0.97 ± 0.001	0.955 ± 0.0006	0.94 ± 0.0006	0.92 ± 0.0015	---
Sucrose (g/l)	0	30	130	230	330	430	530
a_w	1 ± 0	0.999 ± 0.0006	0.995 ± 0.0015	0.987 ± 0.0015	0.98 ± 0	0.973 ± 0.001	0.967 ± 0.001

The values are means of the triplicates plus/minus the standard deviation.

the plugs were left for 2 hours, and extracted as described above. Recovery was determined by comparing the slope of the spiked curve to slope of the curve from diluted samples in methanol-water (3:1). Five blank samples of the 2 strains were also analysed.

Reproducibility of extraction efficiency was further determined from 7 *A. niger* strains (IBT 4983, IBT 18741, IBT 19558, IBT 26774, IBT 28086, IBT 28104, NRRL 567) selected from high, medium and low fumonisin producing strains. Five replicate plates were made from each strain and extracted as described above.

The effect of temperature on the growth and production of fumonisin by *A. niger* and *Fusarium spp*

To assess the effect of temperature on the production of fumonisins, *A. niger* strains were inoculated on CYAS and *Fusarium* strains were inoculated on PDA. The plates were incubated in darkness at 15, 20, 25, 30, 37 or 42 °C respectively for 7 days. The fungi used were the five *A. niger* strains listed in Table 2, and the following *Fusarium* species: *F. napiforme* IBT 2932, *F. nygamai* IBT 8554, *F. verticillioides* IBT 9400, *F. oxysporum* IBT 9514 and *F. proliferatum* IBT 41107.

The effect of glycerol, NaCl and sucrose on the growth and production of fumonisin by *A. niger* and *Fusarium spp*

For investigation of the effect of glycerol, NaCl and sucrose on the production of fumonisin the following experiments were performed: *A. niger* was inoculated on CYA and *Fusarium* on PDA with different concentration of glycerol (0-255 g/l), salt (0-125 g/l) and sucrose (0-530 g/l). For the *Fusarium*-sucrose experiment, the PDA mixture was made from boiled potatoes instead of a commercial blend. In short terms, 200 g potatoes was peeled and diced and autoclaved at 121 °C with 1 liter of water, 15 g agar and 0-530 g sucrose.

The corresponding measured water activity (a_w) values of the media are shown in Table 3. The water activity was

measured with an Aqualab (ADAB Analytical Devices, Stockholm, Sweden). There were no measurable differences in the water activity of the CYA and PDA media. The fungi used were five *A. niger* strains, and the following *F. napiforme* IBT 2932, *F. nygamai* IBT 8554, *F. verticillioides* IBT 9400, *F. oxysporum* IBT 9514 and *F. proliferatum* IBT 41107.

Authors' contributions

JMM, UT and JCF designed the study. JMM performed the experiments, KFN the analytical part and RAS the molecular genetic studies. All authors contributed in drafting the paper and have read and approved the final manuscript.

Additional material

Additional file 1

Effect of temperature on the growth of Aspergillus niger and Fusarium spp. The conidial diameter of 5 Fusarium spp. and 5 Aspergillus niger strains at different temperature in the range of 15-42 °C after 7 days growth.

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Additional file 2

Effect of water activity on the growth of Aspergillus niger and Fusarium spp. The conidial diameter of 5 Fusarium spp. and 5 Aspergillus niger strains at different a_w in the range of 0.92-1 after 7 days growth.

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Acknowledgements

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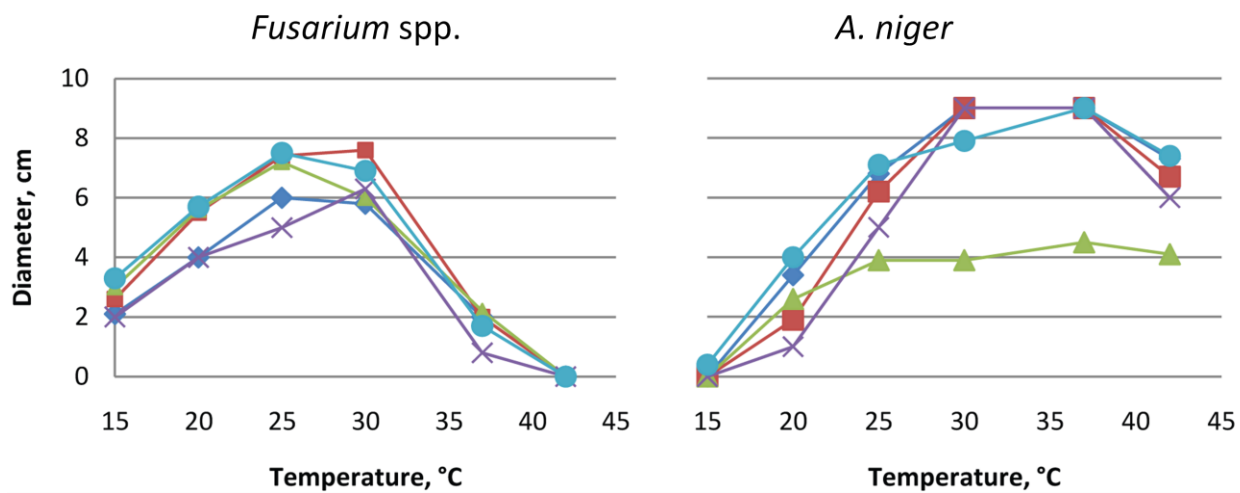


Additional file 1

The conidial diameter of 5 *Fusarium* spp. and 5 *Aspergillus niger* isolates at different temperature in the range of 15-42 °C. All isolates were grown in perforated plastic bags at 25°C for 7 days in darkness. The diameter is an average of the smallest and broadest conidial diameter.

F. napiforme 2932 *F. nygamai* 8554 *F. verticillioides* 9400 *F. oxysporum* 9514
F. proliferatum 41107

A. niger : NRRL 3 NRRL 567 NRRL 2001 IBT 24631 IBT 24634

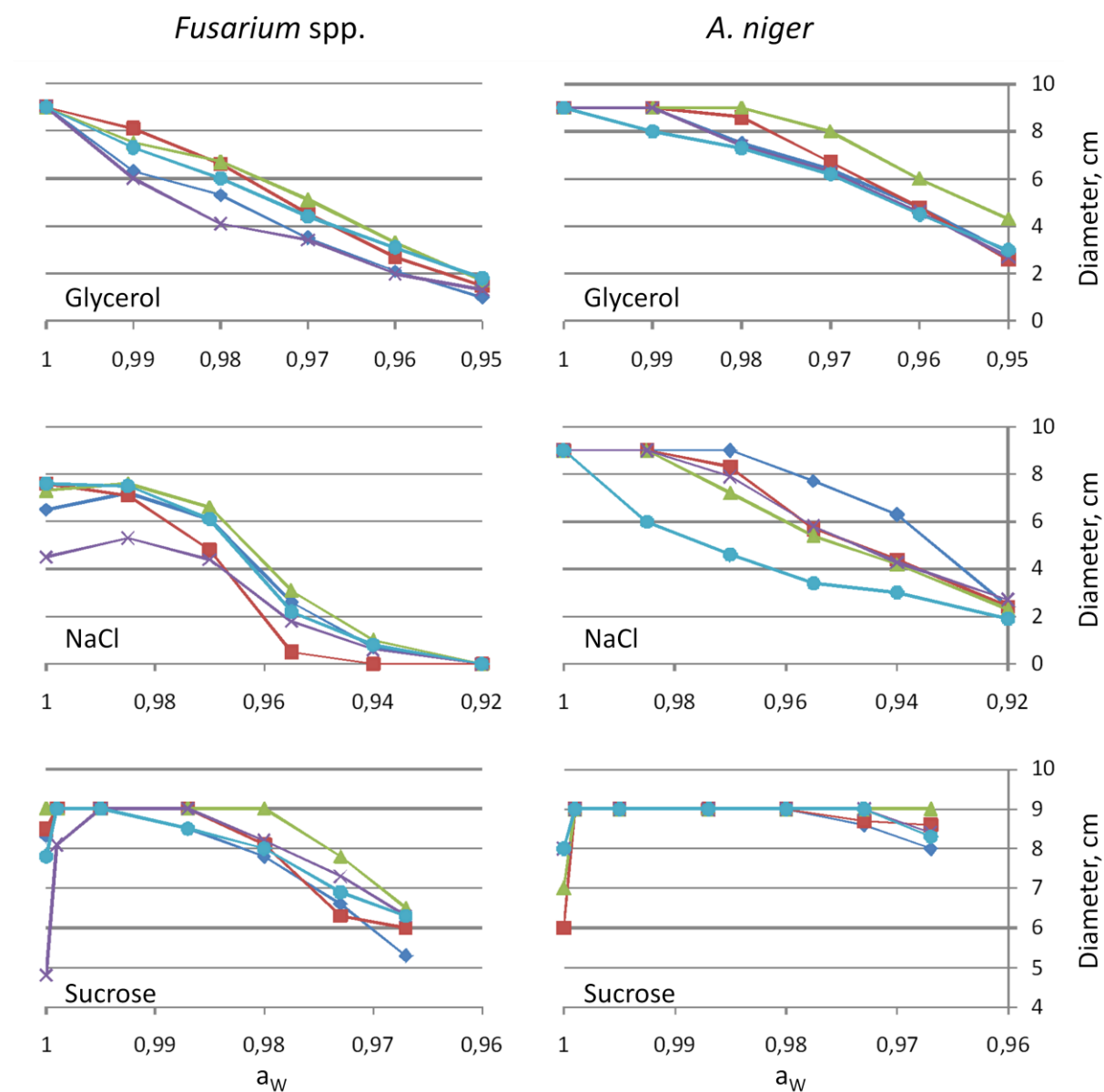


Additional file 2

The conidial diameter of 5 *Fusarium* spp. and 5 *Aspergillus niger* isolates on three different water activity lowering solutes, glycerol, NaCl and sucrose. A_w was in the range of 0.92-1. All isolates were grown in perforated plastic bags at 25°C for 7 days in darkness. The diameter is an average of the smallest and broadest conidial diameter.

—◆— *F. napiforme* 2932 —■— *F. nygamai* 8554 —▲— *F. verticillioides* 9400 —×— *F. oxysporum* 9514
—●— *F. proliferatum* 41107

A. niger : —◆— NRRL 3 —■— NRRL 567 —▲— NRRL 2001 —×— IBT 24631 —●— IBT 24634



Paper 2

” Production of fumonisin B₂ and B₄ by *Aspergillus niger*
on grapes and raisins”

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Production of Fumonisin B₂ and B₄ by *Aspergillus niger* on Grapes and Raisins

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The recent discovery of fumonisin production in *Aspergillus niger*, raises concerns about the presence of these mycotoxins in grapes and raisins as well as other commodities where *A. niger* is a frequent contaminant. Here we investigate the potential production of fumonisins in *A. niger* cultured on grapes and raisins. Sixty-six *A. niger*, 4 *A. tubingensis*, and 16 *A. acidus* strains isolated from raisins were tested for fumonisin production on laboratory media. Neither *A. tubingensis* nor *A. acidus* strains produced fumonisins, but 77% of *A. niger* strains did. None of the strains produced ochratoxin A. Ten selected fumonisin producing *A. niger* strains were further able to produce fumonisin B₂ and fumonisin B₄ on grapes in the range 171–7841 µg fumonisin B₂/kg and 14–1157 µg fumonisin B₄/kg. Four selected strains were able to produce fumonisin B₂ (5–6476 µg/kg) and fumonisin B₄ (12–672 µg/kg) on raisins.

KEYWORDS: Fumonisin; *Aspergillus niger*; mycotoxins; grapes; raisins.

INTRODUCTION

Black Aspergilli are the most common fungi responsible for postharvest decay of fresh fruit (1). Of these *Aspergillus carbonarius* and *A. niger* are very important opportunistic pathogens of grapes causing bunch rot or berryrot and causing raisin mold (2). Since *A. carbonarius* is a much better ochratoxin A producer, the main focus concerning grapes and black Aspergilli has been on *A. carbonarius* and ochratoxin A (3). However, *A. niger* has also been reported to grow and damage a large number of crops and foods worldwide, including corn, peanuts, raisins, onions, mangoes, apples, and dried meat products (1).

A. niger is used for production of single cell protein for feed (4) and is extensively used for production of several organic acids and extracellular enzymes. Furthermore it is used as a transformation host in the biotechnological industry (5, 6). The discovery of putative homologues to the *Fusarium verticillioides* fumonisin gene cluster in three *A. niger* genomes led to the discovery of a fumonisin B₂ production by all three full genome sequenced strains and the ex type culture of *A. niger* (6–8). It has also been shown that *A. niger* is able to produce the biosynthetic precursor fumonisin B₄, which lacks a hydroxyl group on the backbone (9). The structure of fumonisins B₁, B₂, and B₄ are shown in Figure 1.

Fumonisin are important mycotoxins because they are suspected to cause human and animal toxicoses by the consumption of contaminated corn-based food and feeds (10). The fumonisins are structurally similar to sphingolipids and have shown to inhibit the sphingolipid biosynthesis via the ceramide synthase pathway (11). Fumonisin have shown to induce outbreaks of leukoencephalomalacia in horses and pulmonary edema and

hydrothorax in pigs (11, 12). Fumonisin B₁ is hepatocarcinogenic, hepatotoxic, and nephrotoxic in rats and rat liver cancer can be promoted and initiated by fumonisin B₁ (11, 12). Fumonisin B₁ and B₂ have been declared class 2B carcinogens, which are possible human carcinogens (13, 14). The regulatory limit for fumonisins in corn is, according to the U.S. Food and Drug Administration, 2–4 µg/g total fumonisins (15). EEC has a regulatory limit of 0.2–2 µg/g (16). Fumonisin have been isolated from corn and corn-based products (12) such as tortillas (17) and beer (12), as well as rice (12), black tea leaves (18), asparagus (19), and pine nuts (20). Very few factors that affect the fumonisin production by *A. niger* have been investigated; so far only temperature and water activity and a limited amount of media have been investigated (8, 21). Recently fumonisins B₂ and B₄ from *A. niger* were detected in green coffee beans, although the concentrations (up to 10 µg/kg) were well below the regulatory limit for other commodities (9). In a study performed on grape must, 2 out of 12 samples were found to be positive for fumonisin B₂ (0.01 and 0.4 µg/g) (22).

In this study, we investigate the presence of ochratoxin A and fumonisin producing black Aspergilli on raisins, as well as fumonisin production in grapes, dried grapes, and raisins.

MATERIALS AND METHODS

Unless otherwise is stated all solvents were HPLC grade, chemicals were analytical grade, and water was purified on a Milli-Q (Millipore, Billerica, MA). Media were prepared in 9 cm Petri dishes, each with 20 mL of medium. The fumonisin standard was a mixed certified standard containing both fumonisin B₁ and fumonisin B₂ with concentrations of 50.2 and 51.0 mg/L (Biopure, Tulln, Austria). The ochratoxin A standard contained 10 mg/L (Biopure). Fungal strains were three-point inoculated and all samples were incubated in microperforated plastic bags at 25 °C for 7 days in darkness. All water activity measurements were

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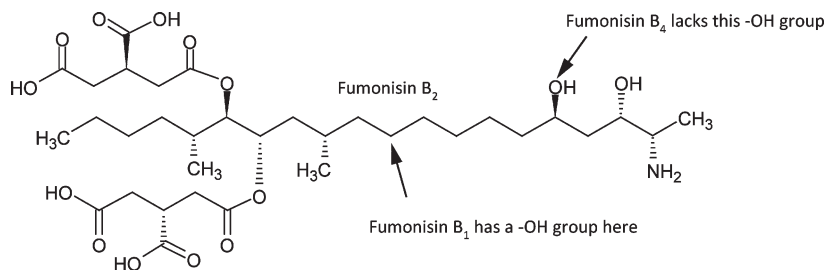


Figure 1. Fumonisin B₂ and the Difference to Fumonisin B₁ and B₄.

performed in triplicate and measured with an Aqualab (ADAB Analytical Devices, Stockholm, Sweden).

Isolation and Characterization of Black Aspergilli Isolated from Raisins. Seventeen different raisin samples were purchased from supermarkets and shops in Denmark, U.S.A., or Uganda. The raisins originated from California, Argentina, or Uganda. Six raisins were selected randomly from each sample, placed on a dichloran 18% glycerol agar (23) plate, and incubated at 25 °C for seven days. All black *Aspergilli* were transferred to Czapek yeast autolysate agar (CYA) and yeast extract sucrose agar (YES) (24) plates and afterward identified using a polyphasic approach (macro- and micromorphology and metabolite profile) according to previous reports (25–27). All strains were subsequently screened for a potential fumonisin production on three different media, CYA, YES, and Czapek yeast autolysate agar with 5% NaCl (CYAS) (24). Fumonisin and ochratoxin A were extracted using an agar plug extraction method (8) with small modifications. Six agar plugs ($D = 6$ mm) were transferred to a 2 mL vial and 800 μ L of methanol/water (3:1) was added. The samples were extracted ultrasonically for 1 h, filtered through a PFTE 0.45 μ m filter (National Scientific, Rockwood, TN), and used directly for LC-TOF-MS analysis. Representative isolates were preserved in the IBT collection (author's address).

Preparation of Grapes. Conidial suspensions for inoculation of grapes were obtained by harvesting spores of each isolate grown on CYA and suspending them in sterile distilled water containing 0.05% of Tween 80 (Merck, Hohenbrunn, Germany) and 0.05% agar (Bie & Berntsen, Rødovre, Denmark). The final concentration of the conidia was assessed by using a counting chamber and was adjusted to 10^6 conidia/mL.

Fresh grapes (Thompson seedless), cut in half, were placed in an empty Petri dish and inoculated with 10 μ L of the *A. niger* conidia suspensions. The grape experiment was prepared in duplicate. Ten strains randomly selected from the before mentioned screening were selected for the grape experiment, IBT 28747, 28753, 28934, 28937, 28948, 28964, 28965, 28966, 28994, and 29019.

Preparation of Commercial Raisins. From one brand, two subsamples of 125 g raisins (California, Thompson seedless) were randomly selected and surface sterilized with hypochlorite (3%) for 15 min and then washed twice with sterile water (weight gain 7%). One of the two subsamples was covered with water and both subsamples were kept at 5 °C for 24 h (weight gain 50%). The raisins placed in water were transferred to an empty Petri dish; the other subsample was placed on a water agar separated by a filter (pore size 1 μ m). Four *A. niger* strains with the highest production of fumonisins in grapes were chosen; these were IBT 28753, 28934, 28948, and 29019. The raisins were inoculated with 10 μ L of the *A. niger* conidia suspensions in quadruplicates as described above.

Preparation of Dried Grapes. Portions of 300 g grapes (size, 2–3 g) (Thompson seedless) were taken at random and dried in an oven at 75 °C for 4 h (weight loss, 32%, initial a_w 0.76 ± 0.04), 5 h (weight loss, 38%, initial a_w 0.69 ± 0.01), and 6.5 h (weight loss, 52%, initial a_w 0.54 ± 0.09). Subsequently, they were placed in an empty Petri dish and inoculated with *A. niger* as described above. All samples were as a minimum prepared in triplicates. Four *A. niger* strains with the highest production of fumonisins in grapes were chosen; these were IBT 28753, 28934, 28948, and 29019.

Extraction of Grapes and Raisins for Chemical Analysis. One grape or raisin was weighed and placed in a 5 mL cryo tube with 5–10 steel balls ($D = 3$ mm), 5 mL methanol/water (3:1) was added. The cryo tubes were shaken in a Mini Beadbeater 96 (Biospec Product Inc., Bartlesville, OK) for 5 min. The whole mixture was transferred to a 15 mL Falcon tube.

Afterward it was placed on a shaking table (at an angle of 45°) for 60 min at 150 rpm. The Falcon tubes were centrifuged at 8000 g for 4 min and 200 μ L extract was transferred to a 2 mL HPLC vial and used directly for LC-MS/MS analysis. Because of nonavailability of fumonisin B₄ as analytical standard, external quantification was performed with fumonisin B₂ and similar response factors were assumed for the two mycotoxins.

LC-TOF-MS Conditions. The screening for a fumonisin in culture extracts production was performed using a LC system coupled to an orthogonal TOF mass spectrometer (Micromass LCT, Manchester, U.K.) equipped with an electrospray source (25). The column used was a 50 \times 2 mm i.d., 3 μ m Gemini C₆-phenyl column (Phenomenex, Torrance, CA) with a linear gradient starting from 30% acetonitrile in water (both 20 mM formic acid) to 60% acetonitrile for 5 min at a flow rate of 300 μ L/min, which was then increased to 100% acetonitrile in 1 min and a flow of 0.5 mL/min, keeping this for 3.5 min before returning to the start conditions in 6 min. The mass spectrometry was performed in ESI⁺.

LC-MS/MS Conditions. LC-MS/MS analysis was performed as previously described (9) with an Agilent HP 1100 liquid chromatography system (Waldbronn, Germany) coupled to a Quattro Ultima triple mass spectrometer (Micromass, Manchester, UK) with ESI source. The gradient was as described before but with the gradient starting at 20% CH₃CN and going to 55% CH₃CN in 5 min and then to 100% in 30 s. Tandem mass spectrometry was performed in ESI⁺ at a source flow at 700 L/h nitrogen at 350 °C. Nitrogen was also used as collision gas, and the MS operated in multiple reaction monitoring mode at the following transitions: fumonisin B₂ quantifier m/z 706 \rightarrow 336 cone 50 V, collision 40 V, dwell time 50 ms, qualifier m/z 706 \rightarrow 512, cone 50 V, collision 25 V, dwell time 100 ms (ion ratio, 1.7–2.1 (transition 1/2)); fumonisin B₄ quantifier m/z 690 \rightarrow 320 cone 50 V, collision 35 V, dwell time 50 ms, qualifier m/z 690 \rightarrow 514 a, cone 50 V, collision 30 V, dwell time 100 ms (ion ratio, 1.4–1.8 (transition 1/2)); ochratoxin A quantifier m/z 404 \rightarrow 239, cone 30 V, collision 33 V, dwell time 100 ms, qualifier m/z 404 \rightarrow 358, cone 30 V, collision 25 V, dwell time 100 ms (ion ratio, 1.4–1.7 (transition 1/2)).

Validation. The method was validated by spiking raisins and grapes with 100 μ L of a mixture of fumonisin B₂ and ochratoxin A standards to the following six different final levels: fumonisin B₂, 15, 25, 50, 150, 200, and 300 μ g/kg; ochratoxin A, 3, 5, 10, 30, 40, and 60 μ g/kg in triplicate. The samples were left to dry for 2 h prior to extraction. Both the raisins and the grapes were extracted as described above.

RESULTS AND DISCUSSION

Preliminary testing (results not shown) of strong anion exchange purification (SAX) showed low or no recoveries of fumonisins from the grapes, presumably due to competition from high amounts organic acids from the grapes and/or from *A. niger* strains themselves. Furthermore immunoaffinity purification (results not shown) yielded viscous brown extracts from raisin extracts and was thus not suitable either. Consequently, it was decided to analyze the crude extracts.

For validation, standard curves were prepared and shown to be linear. On grapes, the relative standard deviation of the lowest point (15 μ g/kg) was 18% (fumonisin B₂ level); for raisins it was 7%. All R^2 , level of detection (LOD), and level of quantification (LOQ) values are shown in Table 1. Because of the high fumonisin concentrations detected in the strains, LOD_{fum} were not

experimentally addressed, but estimation from the lowest point down to s/n indicates LOD of 5 $\mu\text{g/kg}$. Control analysis of conidial suspensions shows that in approximately 10 000 spores the concentration of fumonisin was in picogram levels and would be diluted 1:500 in the final extract thus not influencing the result.

The screening resulted in isolation of 86 black *Aspergillus* strains; of these 66 were *A. niger*, 4 *A. tubingensis*, and 16 *A.*

Table 1. R^2 , Level of Determination (LOD), and Level of Quantification (LOQ) Determined from Spiked Raisin and Grape Samples

matrix	fumonisin B ₂			ochratoxin A		
	R^2 (n = 18)	LOD ^a	LOQ	R^2 (n = 18)	LOD ^b	LOQ
grape	0.993	5 $\mu\text{g/kg}$	25 $\mu\text{g/kg}$	0.96	3 $\mu\text{g/kg}$	10 $\mu\text{g/kg}$
raisin	0.990	3 $\mu\text{g/kg}$	25 $\mu\text{g/kg}$			

^a Estimated from the lowest point (15 $\mu\text{g/kg}$) to s/n (1:5). ^b The s/n is 1:5.

Table 2. Production of Fumonisins by *Aspergillus niger* in Grapes after 7 Days Growth at 25°C

isolate	fumonisin B ₂ ($\mu\text{g/kg}$)		fumonisin B ₄ ($\mu\text{g/kg}$) ^a		ochratoxin A $\mu\text{g/kg}$ ^b
	sample 1	sample 2	sample 1	sample 2	
IBT					
28747	951	1283	75	189	nd
28753	7703	7979	772	1542	nd
28934	2098	5376	158	612	nd
28937	1045	1941	253	265	nd
28948	3380	4772	349	509	nd
28964	91	251	5	23	nd
28965	1074	2414	148	344	nd
28966	1261	5123	220	634	nd
28994	432	2422	36	158	nd
29019	4303	11189	143	405	nd

^a Assuming same response factor as fumonisin B₂. LOD: 5 $\mu\text{g/kg}$. ^b LOD: 3 $\mu\text{g/kg}$. nd: not detected.

acidus (= *A. foetidus* var. *acidus* (28)). None of the *A. tubingensis* and *A. acidus* strains produced fumonisins, but 77% of the *A. niger* produced fumonisin B₂ (LC-TOF). Similar results were found among a limited number of grape derived strains (22). A similar recent study performed on green coffee beans found the a similar fraction (76%) of *A. niger* as fumonisin B₂ producers (9). The percentage of ochratoxin A -producing *A. niger* is most often reported in the interval 0–30% (3, 29–31), but in one case a percentage as high as 41% has been reported (32). The percentages of ochratoxigenic strains of *A. niger* are very low compared to our 77% fumonisin producing *A. niger*. Therefore this fumonisin production by *A. niger* could present an even larger food safety problem.

The 10 *A. niger* strains with the highest measured fumonisin production were selected for further experiments, IBT 28747, 28753, 28934, 28937, 28948, 28964, 28965, 28966, 28994, and 29019. All 10 strains were able to produce fumonisins in grapes (Table 2). Fumonisin B₂ was produced as the major component, followed by fumonisin B₄. The mean production of fumonisin B₂ varied almost 50 fold from 171 $\mu\text{g/kg}$ (IBT 28964) to 7841 $\mu\text{g/kg}$ (IBT 28753) and fumonisin B₄ varied almost 70 fold from 14 $\mu\text{g/kg}$ (IBT 28964) to 1157 $\mu\text{g/kg}$ (IBT 28753). Selected transitions from the LC-MS/MS analysis of infected and spiked grapes are shown in Figure 2. The measured fumonisin concentrations are similar, compared to those previously described by Logrieco et al. (200–2500 $\mu\text{g/kg}$) (22).

The four strains with the highest production of fumonisins on grapes were selected for the work with dried grapes and raisins. On the dried grapes, fumonisin B₂ and fumonisin B₄ were also produced (Table 3). The concentration of fumonisin B₂ was in the range of 91–1747 $\mu\text{g/kg}$ and fumonisin B₄ 9–69 $\mu\text{g/kg}$. The fumonisin production was highest in the grapes with the lowest weight losses and the lowest production was in all coincidences found in the grapes with the lowest water content.

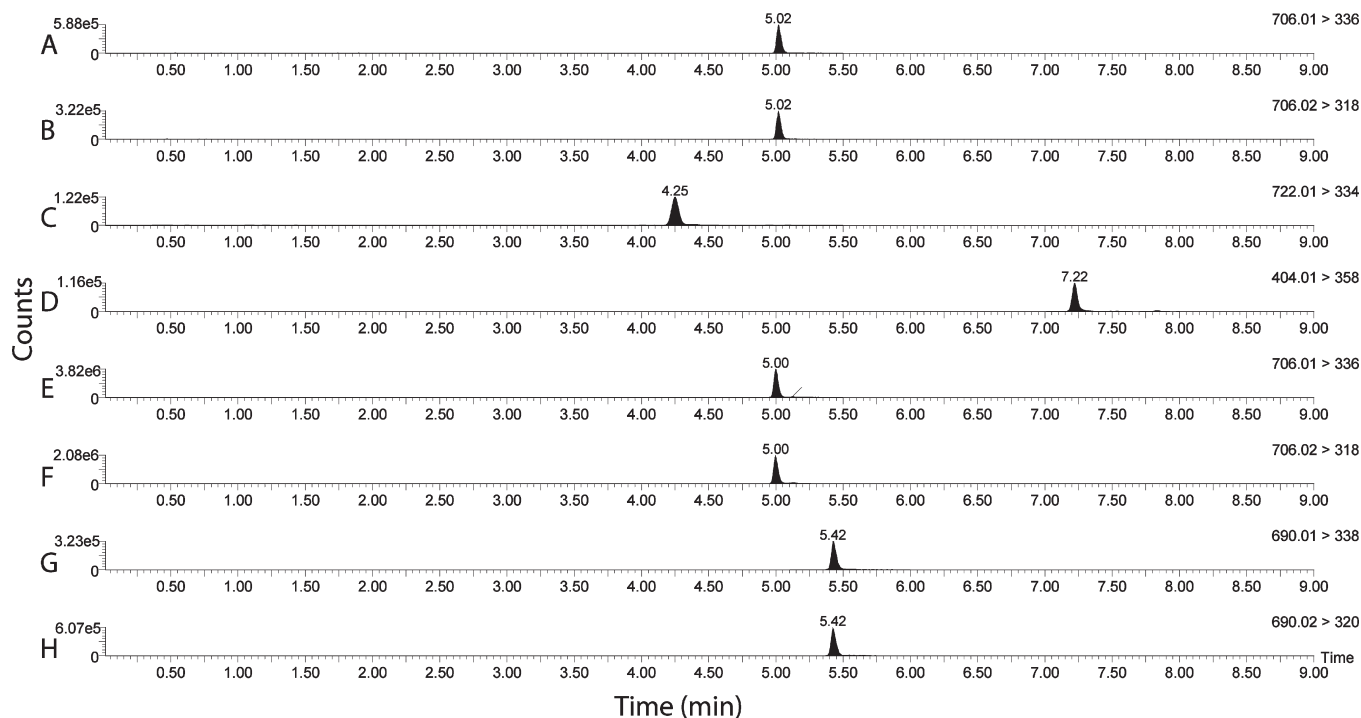


Figure 2. Results from the LC-MS/MS analysis. Transitions of Ochratoxin A, Fumonisin B₁, B₂ and B₄ in Spiked Grape and Infected Grape. (A) Quantifier of Fumonisin B₂ from a Spiked Grape (150 $\mu\text{g/kg}$). (B) Qualifier of Fumonisin B₂ from a Spiked Grape (150 $\mu\text{g/kg}$). (C) Quantifier of Fumonisin B₁ from a Spiked Grape (150 $\mu\text{g/kg}$). (D) Quantifier of Ochratoxin A from a Spiked Grape (30 $\mu\text{g/kg}$). (E) Quantifier of Fumonisin B₂ from an Infected Grape (7841 $\mu\text{g/kg}$ IBT 28753). (F) Qualifier of Fumonisin B₂ from an Infected Grape (7841 $\mu\text{g/kg}$ IBT 28753). (G) Quantifier of Fumonisin B₄ from an Infected Grape (1157 $\mu\text{g/kg}$ IBT 28753). (H) Qualifier of Fumonisin B₄ from an Infected Grape (1157 $\mu\text{g/kg}$ IBT 28753).

Table 3. Production of Fumonisin by *Aspergillus niger* in Dried Grapes after 7 Days Growth at 25°C

isolate IBT	weight loss %	initial a_w	fumonisin B ₂ $\mu\text{g/kg}^a$	fumonisin B ₄ $\mu\text{g/kg}^{a,b}$	ochratoxin A $\mu\text{g/kg}^c$
28753	32	0.76 \pm 0.04	1747 \pm 316	48 \pm 22	nd
	38	0.69 \pm 0.01	1128 \pm 329	48 \pm 16	nd
	52	0.54 \pm 0.09	91 \pm 73	9 \pm 4	nd
28934	32	0.76 \pm 0.04	1363 \pm 954	69 \pm 51	nd
	38	0.69 \pm 0.01	1033 \pm 460	50 \pm 23	nd
	52	0.54 \pm 0.09	636 \pm 397	24 \pm 16	nd
28948	32	0.76 \pm 0.04	313 \pm 323	54 \pm 47	nd
	38	0.69 \pm 0.01	154 \pm 71	27 \pm 11	nd
	52	0.54 \pm 0.09	144 \pm 104	29 \pm 19	nd
29019	32	0.76 \pm 0.04	203 \pm 106	25 \pm 11	nd
	38	0.69 \pm 0.01	214 \pm 137	28 \pm 13	nd
	52	0.54 \pm 0.09	102 \pm 76	10 \pm 5	nd

^a The values are means of the quadruplicates, plus/minus the standard deviation. ^b Assuming same response factor as Fumonisin B₂. LOD: 5 $\mu\text{g/kg}$. ^c LOD: 3 $\mu\text{g/kg}$. nd: not detected.

Table 4. Production of Fumonisin by *Aspergillus niger* in Raisins after 7 Days Growth at 25°C

isolate IBT	weight gain	initial a_w	fumonisin B ₂ $\mu\text{g/kg}^a$	fumonisin B ₄ $\mu\text{g/kg}^{a,b}$	ochratoxin A $\mu\text{g/kg}^c$
28753	7% (agar)	0.77 \pm 0.05	1160 \pm 686	105 \pm 56	nd
	50% (water)	0.95 \pm 0.001	407 \pm 249	47 \pm 25	nd
28934	7% (agar)	0.77 \pm 0.05	229 \pm 58	27 \pm 7	nd
	50% (water)	0.95 \pm 0.001	5 \pm 5	nd	nd
28948	7% (agar)	0.77 \pm 0.05	459 \pm 430	45 \pm 31	nd
	50% (water)	0.95 \pm 0.001	112 \pm 51	12 \pm 4	nd
29019	7% (agar)	0.77 \pm 0.05	6476 \pm 1139	356 \pm 46	nd
	50% (water)	0.95 \pm 0.001	784 \pm 636	672 \pm 422	nd

^a The values are means of the quadruplicates, plus/minus the standard deviation. ^b Assuming same response factor as Fumonisin B₂. LOD: 5 $\mu\text{g/kg}$. ^c LOD: 3 $\mu\text{g/kg}$. nd: not detected.

Two experiments were carried out with the raisins, one where the water activity (a_w) initially was low and over time increased toward 1, and another where the water activity started high and decreased over time. All of the tested *A. niger* strains were found to produce fumonisin B₂ and fumonisin B₄ on both types of raisins (Table 4). In all cases the production of fumonisins was highest when the raisins were kept moist. Fumonisin B₂ and fumonisin B₄ were produced in the range of 229–6476 and 27–356 $\mu\text{g/kg}$. The raisins with a decreasing water activity had a fumonisin B₂ concentration of 5–784 $\mu\text{g/kg}$ and fumonisin B₄ of 12–672 $\mu\text{g/kg}$.

In the dried grapes the production of fumonisins decreased as the water activity was lowered. This correlates with earlier results that the lower a_w , the more reduced fumonisin production by *A. niger* (21). The reason why the raisins with the lowest a_w supported fumonisin production could be because the wet grapes dried quickly and the raisins on the water agar were kept moist with a high a_w on the surface. For the four tested strains, in the three experiments all had the highest production of fumonisins in grapes.

This is the first report describing the production of fumonisins B₂ and B₄ by *A. niger* in dried grapes and raisins. All the *A. niger* produced more fumonisin B₂ than fumonisin B₄. No production of ochratoxin A (LOD: 3 $\mu\text{g/kg}$) was detected in either the grape or raisin experiments. This is supported by earlier studies that have shown *A. carbonarius* and not *A. niger* as the most important source of ochratoxin A in raisins (29, 33, 34).

The presence of a fumonisin production in grapes and a detectable content of fumonisin in grape must (22) indicates that fumonisins may be found in wine unless degraded during the fermentation or storage. Consequently, a larger number of wines as well as grape musts need to be investigated to track the fate of fumonisins in all production steps from grapes to wine.

Fumonisin produced by *A. niger* may be an overlooked health risk in a broad variety of foods, because *A. niger* is a common

food spoilage organism (1). New knowledge within this field is important, particularly because corn and derived products are currently the only major products that are monitored for fumonisins.

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Paper 3

” Widespread occurrence of the mycotoxin Fumonisin B₂ in wine”

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Fumonisin is an important mycotoxin because they are suspected to cause human and animal toxicoses by the consumption of contaminated corn-based food and feeds. However, with the discovery of fumonisin production in grapes by *Aspergillus niger*, wine may also be a fumonisin-containing commodity. In the present study, we have developed a simple and quantitative cation-exchange-based purification method for the subsequent isotope dilution liquid chromatography–tandem mass spectrometry (LC–MS/MS) determination of fumonisins in wine. A comparative study of seven different solid-phase extraction (SPE) columns showed that polymeric mixed-mode reversed-phase (RP) cation-exchange columns were superior to classic silica-based cation and mixed-mode cation-exchange columns. A total of 77 wine samples from 13 countries were subsequently tested, and surprisingly, 18 (23%) were found to contain fumonisin B₂ in the range of 1–25 µg/L. These findings were further confirmed by immunoaffinity purification and re-analysis of the positive cation-exchanged extracts.

KEYWORDS: Fumonisin; wine; grapes; *Aspergillus niger*

INTRODUCTION

Black aspergilli are some of the most common fungi responsible for postharvest decay of fresh fruit (1). Of these are *Aspergillus carbonarius* and *Aspergillus niger*, very important opportunistic pathogens of grapes causing bunch rot or berryrot (2). Because *A. carbonarius* can produce large amounts of ochratoxin A, the main focus concerning grapes, wine, and black aspergilli has therefore been on this fungus and ochratoxin A (3–6). The recent discovery of fumonisins B₂ (7), B₄ (8), and B₆ (9) (Figure 1) from *A. niger* and the production of fumonisins by *A. niger* in coffee and especially grapes, raisins, and must (8, 10, 11) have raised concerns of the possible general presence of fumonisins in wine.

Fumonisin is an important mycotoxin because they are suspected to cause human and animal toxicoses by the consumption of contaminated corn-based food and feeds (12). To avoid possible detrimental health effects, the United States Food and Drug Administration (U.S. FDA) recommends that maize should not be used for human consumption when contaminated with more than 2–4 mg/kg total fumonisins (13), whereas the European Union (EU) has a regulatory limit of 0.2–2 mg/kg (14).

Thus far, no detection method for fumonisins in wine has been described. Non-purified but highly concentrated extracts of must, grapes, and raisins have been analyzed using liquid chromatography–tandem mass spectrometry (LC–MS/MS) (10, 11). However, for low concentrations in a high complexity of this product, wine (e.g., containing polyphenolics, flavonoids, tannins, and

anthocyanins), this approach will clearly limit sensitivity and poses ion-suppression problems unless a much more sensitive instrument is used and, thus, less sample can be injected. We therefore decided to develop a fast solid-phase extraction method with subsequent LC–MS/MS analysis for the determination of fumonisins in wine. The method described here is to our knowledge the first quantitative method for fumonisins using cation exchange.

MATERIALS AND METHODS

Unless otherwise stated, all solvents were high-performance liquid chromatography (HPLC)-grade and other chemicals were analytical-grade and from Sigma-Aldrich (St. Louis, MO). Water was purified on a Milli-Q system (Millipore, Bedford, MA). The fumonisin standard was a mixed certified standard containing both fumonisins B₁ and B₂, with concentrations of 50.2 and 51.0 mg/L (Biopure, Tulln, Austria). Relevant samples were added U-¹³C₃₄-fumonisin B₂ (10.52 µg/mL, Biopure) to a total concentration of 10 µg/L.

Sample Preparation. Wine samples of 10 mL (spiked or non-spiked, pH~3.5–4) were, unless otherwise stated, passed through a 30 mg/3 mL Strata X-C column (Phenomenex, Torrance, CA), which was previously conditioned with 1 mL of methanol and 1 mL of water. To further acidify the column, 1 mL of water with 2% formic acid was added. Afterward, the column was washed with 1 mL of methanol and the fumonisins were eluted with methanol/water (1:4) with 2% NH₄OH water into a 2 mL HPLC vial and used directly for analysis.

The following other cation-exchange columns were tested with the above method as well as small variations of this: Oasis MCX, 30 mg (Waters, Milford, MA) and the silica based ones, HCB, 130 mg (C₈, SCX, Biotage, Uppsala, Sweden); HCB-3, 130 mg (C₁₈, SCX, Biotage); HCB-5, 130 mg (C₄, SCX, Biotage); and SCX, 100 mg (Phenomenex).

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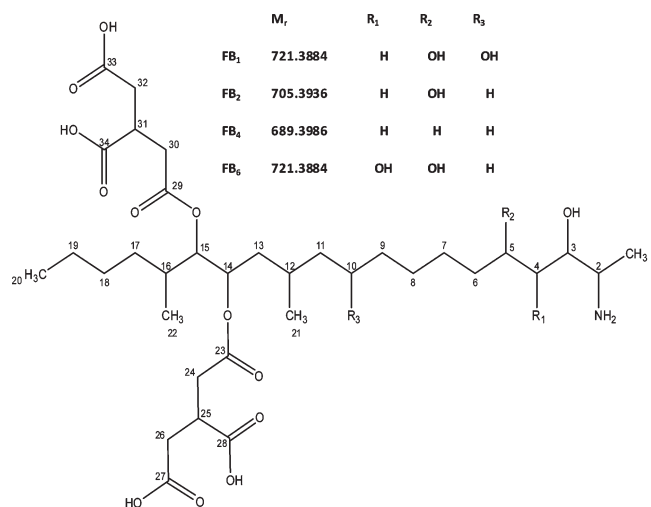


Figure 1. Structure of fumonisin B₂ and a comparison to fumonisins B₁, B₄, and B₆.

SAX purification was also tested on 100 mg columns (Phenomenex). The following procedure was used: The pH of the wine was adjusted to 7.0 ± 0.2 with 1M NaOH, and the columns were conditioned with 1 mL of methanol and 1 mL of water, followed by the addition of wine. The cartridge was washed with 1 mL of methanol/water (7:3) and 1 mL of methanol. The samples were eluted with methanol with 2% formic acid and used directly for analysis. Each sample was spiked with an internal standard of U-¹³C₃₄-fumonisin B₂ to a total concentration of 10 μ g/L.

Validation. The fumonisin B₂ extraction was validated by spiking portions of 10 mL of wine to the following concentrations of fumonisin B₂: 0 (blank), 2, 5, 15, 50, 150, and 450 μ g/L. The experiments were performed on different days, with all levels in triplicates each day. The apparent recovery was calculated from methanol/water (20:80) diluted samples with the following concentrations of fumonisin B₂: 2, 5, 15, 50, 150, and 450 μ g/L. An internal standard of U-¹³C₃₄-fumonisin B₂ was added to each sample to a total volume of 10 μ g/L. To determine the apparent recovery, the slope of spiked wine samples and MeOH standards was compared for similar concentrations.

Wine Samples. Wine samples for screening were either bought in a local supermarket or kindly provided by employees at the Department of Systems Biology, Technical University of Denmark. In total, 77 wines from 13 countries were tested, including red ($n = 56$), white ($n = 13$), port ($n = 6$), champagne ($n = 1$), rosé ($n = 1$), and madeira ($n = 1$) wines. The wines were from the years 1991–2009 and from Argentina ($n = 1$), Australia ($n = 6$), Chile ($n = 3$), China ($n = 1$), France ($n = 22$), Germany ($n = 1$), Italy ($n = 23$), New Zealand ($n = 1$), Portugal ($n = 7$), Romania ($n = 1$), Spain ($n = 9$), South Africa ($n = 2$), and the U.S. (California) ($n = 1$). All information on the wines was taken from the label of the bottle.

LC–MS/MS Conditions. LC–MS/MS analysis was performed as previously described (11) but with minor changes. Briefly described, the LC–MS/MS analysis was performed on a Quattro Ultima triple mass spectrometer (Micromass, Manchester, U.K.) with a z-spray electrospray ionization (ESI) source. Separations were performed on a 50 \times 2 mm inner diameter, 3 μ m Gemini C₆-phenyl column (Phenomenex, Torrance, CA). Using a linear gradient starting from 20% acetonitrile in water (both supplemented with 20 mM formic acid) to 55% acetonitrile for 6 min at a flow rate of 0.3 mL/min, which was then increased to 100% acetonitrile in 30 s at a flow of 0.5 mL/min, keeping this for 3.5 min before returning to the start conditions in 6 min. MS/MS was performed in ESI⁺, and the MS operated in multi reaction monitoring (MRM) mode at the following transitions: fumonisin B₂, quantifier m/z 706 \rightarrow 336, cone, 50 V; collision, 40 V; dwell time, 50 ms; qualifier m/z 706 \rightarrow 512, cone, 50 V; collision, 25 V; dwell time, 100 ms; fumonisin B₄, quantifier m/z 690 \rightarrow 320, cone, 50 V; collision, 35 V; dwell time, 50 ms; qualifier m/z 690 \rightarrow 514, cone, 50 V; collision, 30 V; dwell time, 100 ms; fumonisins B₁ and B₆, quantifier m/z 722 \rightarrow 334, cone, 50 V; collision, 40 V; dwell time, 50 ms; qualifier m/z 722 \rightarrow 528, cone, 50 V; collision, 25 V; dwell time, 100 ms; and U-¹³C₃₄-fumonisin B₂ m/z 740 \rightarrow 358, cone, 50 V; collision, 50 V; dwell time, 50 ms.

Verification of Positive Samples. Because a significant number of the positive samples were close to the limit of quantitation (LOQ), it was decided to verify the findings by purification on FumoniTest immunoaffinity columns (Vicam, Watertown, MA). Half of the purified extract (500 μ L) from the LC–MS/MS screening was diluted with phosphate-buffered saline (PBS) buffer to a final volume of 10 mL. The pH was adjusted to 7 ± 0.3 with 1 M HCl before it was loaded onto the immunoaffinity column (1–2 drops/s). The column was washed with 10 mL of PBS buffer; air was pushed through to remove buffer; and possible fumonisins were eluted with 1.5 mL of methanol (1 drop/s). The eluate was evaporated to dryness under a stream of nitrogen at approximately 45 °C, and the dried residue was redissolved in 100 μ L of methanol/water (1:4) and re-analyzed by LC–MS/MS (5 μ L injected).

RESULTS AND DISCUSSION

Sample Purification. Fumonisin have mainly been purified by anion-exchange or immunoaffinity columns (15, 16). Our first approach was the anion-exchange approach using SAX columns, which could be used to target not only fumonisins but also ochratoxin A. However, SAX purification was irreproducible and suffered from poor recovery ($\ll 50\%$), as also observed in a previous study on the analysis of fumonisins from grapes, presumably because of the competition from organic acids from the grapes (11). Subsequently, it was decided to use cation exchange, because fumonisins have an amine group, thereby avoiding the anionic compounds. In our first attempt, we compared Strata SCX (silica-based) to polymeric Strata X-C, with the latter being superior, because of stable flow and better recovery. To find an optimal column, we further compared five different mixed-mode reversed-phase (RP) cation-exchange SPE columns for their abilities to extract fumonisins from wine. The worst tested was clearly the anion-exchange column (SAX) with only 11% extracted fumonisin compared to the best column (MCX). Concerning the C₄, C₈, and C₁₈ mixed-mode SCX columns, there was a recovery of 71, 50, and 12%, respectively, indicating that an increase in the hydrophobicity of the column resulted in a more inefficient recovery of fumonisins. Two of the six tested columns, Oasis MCX and Strata X-C, clearly gave recoveries of 86 and 83%, respectively, with the latter being slightly more reproducible.

To determine the loading capacity of fumonisins on the Strata X-C, seven concentrations within the range of 100–1500 μ g/L fumonisin spiked wine were loaded onto columns and, after LC–MS/MS analysis, the eluted fumonisin showed a linear relationship with no signs of reaching a maximal capacity, indicating a maximal capacity above 1500 μ g/L.

Validation. The three calibration curves, one from each day, from spiked samples, had the following R^2 values: 0.9928, 0.9957, and 0.9978, with a relative standard deviation (RSD) of the slope being 10%. The RSD of the spiked samples ($N = 9$) varied within (0.6–35%), with an average RSD of 15%. The lowest concentration (2 μ g/L) gave a s/n of 20:1, indicating that the limit of detection (LOD) was approximately 4 times lower (assuming a s/n of 5:1). Recovery was found to be 60–93%, with an average of 83% and an average RSD of 12%.

Screening of Wine Samples. Analysis of the 77 wine samples demonstrated the occurrence of fumonisin B₂ in 23% of the samples (18 of 77). This method gives four identification points via retention time and two MRMs and, thus, exceeding the requirements for identification (three points) (17). Because this is the first report describing fumonisin B₂ in wine and that a high number of samples were close to the LOQ, it was decided to further confirm the results by immunoaffinity purification. This was not formally validated because the use of the ¹³C standard validates each purification. By comparison of the peak areas of

Table 1. Overview of the Tested Wine Samples and Their Fumonisin Content^a

sample number	country	grape sort	year	wine type	fumonisin ($\mu\text{g/L}$)
1	Australia	Chardonnay	2008	white	nd ^b
2	Spain	Monastrell		red	2.1
3	Australia	Shiraz, Cabernet Sauvignon	2006	red	nd
4	Italy			red	nd
5	South Africa	Cabernet Sauvignon	2007	red	2.5
6	Australia	Shiraz		rose	nd
7	Romania	Pinot Noir	2006	red	20
8	Spain	Tempranillo Garnacha	2007	red	nd
9	Italy	Merlot, Corvina	2007	red	7.0
10	Australia	Shiraz	2008	red	1.0
14	Italy	Corvina, Rondinella, Molinarea, Negrana	2007	red	6.7
17	France	Sauvignon Blanc, Semillon Muscadelle	2003	white	nd
18	Italy	Corvina, Rondinella	2007	red	nd
19	France	Chardonnay	2008	white	nd
20	France		2007	champagne	nd
21	Chile	Chardonnay	2008	white	nd
24	France	Carignan, Grenache, Syrah	2003	red	5.1
25	Italy		2005	red	nd
26	France	Chardonnay	2008	white	nd
27	Spain	Tempranillo, Gaunacha	2005	red	1.9
29	Italy	Shiraz	2007	red	nd
30	Italy		2003	red	nd
31	Italy		1998	red	nd
33	California, U.S.A.	Zinfandel	1998	red	25
34	Portugal		2007	port	nd
36	Italy	Grecanico, Chardonnay	2008	white	nd
37	France	Grenache, Syrah	2003	red	1.3
38	Portugal		2002	madeira	nd
39	Italy	Montepulciano, Aglianico	2005	red	2.1
40	Italy	Merlot, Corbina	2007	red	3.9
42	Chile	Chardonnay	2008	white	nd
43	Australia		2008	white	nd
44	South Africa		2008	white	nd
45	France	Grenache, Syrah	2007	red	nd
46	Italy	Malvasia, Trebbiano	2008	white	nd
47	Italy	Montalcino, Sangiovese Grosso, Brunello	2000	red	nd
49	Italy	Nebbiolo	1996	red	nd
50	Australia	Shiraz	2007	red	nd
51	China	Cabernet Sauvignon	2007	red	nd
52	Germany	Pinot Noir	2004	red	nd
53	France	Cabernet Sauvignon, Cabernet Franc, Merlot, Petit Verdot	1999	red	nd
54	France		1998	red	nd
57	New Zealand	Sauvignon Blanc	2008	white	nd
58	Italy	Negroamaro, Primitivo	2006	red	1.6
59	France		2003	red	nd
63	Italy	Primitivo di Manduria	2007	red	1.2
64	Italy		2005	red	nd
65	France		2007	red	nd
66	Italy	Corvina, Rondinello, Molinara	2004	red	3.2
67	Spain	Monastrell	2008	red	nd
68	France	Grenache, Syrah, Mourvedre	2008	red	6.5
71	France	Merlot, Cabernet Sauvignon, Cabernet Franc	2003	red	nd
72	France		2002	red	nd
73	Portugal			port	nd
74	France			white	nd
78	Italy	Cabernet Sauvignon, Cabernet Franc	2001	red	nd
79	France	Riesling	2008	white	nd
80	Spain		2004	red	nd
84	France	Cabernet Sauvignon	1996	red	nd
89	Italy		2008	white	1.0
90	France		2007	red	nd
91	Italy	Sangiovese, Rondinella, Corvina	2006	red	nd
92	Spain	Tempranillo	2005	red	nd
93	France		1991	red	nd
94	France		1996	red	nd
95	Italy			red	nd
96	Chile	Carmenere	2008	red	nd

Table 1. Continued

sample number	country	grape sort	year	wine type	fumonisin ($\mu\text{g/L}$)
99	Argentina	Cabernet, Merlot	2008	red	nd
100	Italy			rose	nd
101	Spain			red	nd
102	France			red	nd
103	Portugal			port	nd
104	Portugal			port	2.8
105	Portugal	Cabernet Sauvignon, Merlot Noir, Cabernet Franc	2003	port	nd
106	Spain			red	nd
107	Portugal			red	nd
108	France			red	nd

^aThe description of the wines was read off the label of the bottle. Blank cells indicate that information was not available. ^bnd = not detected.

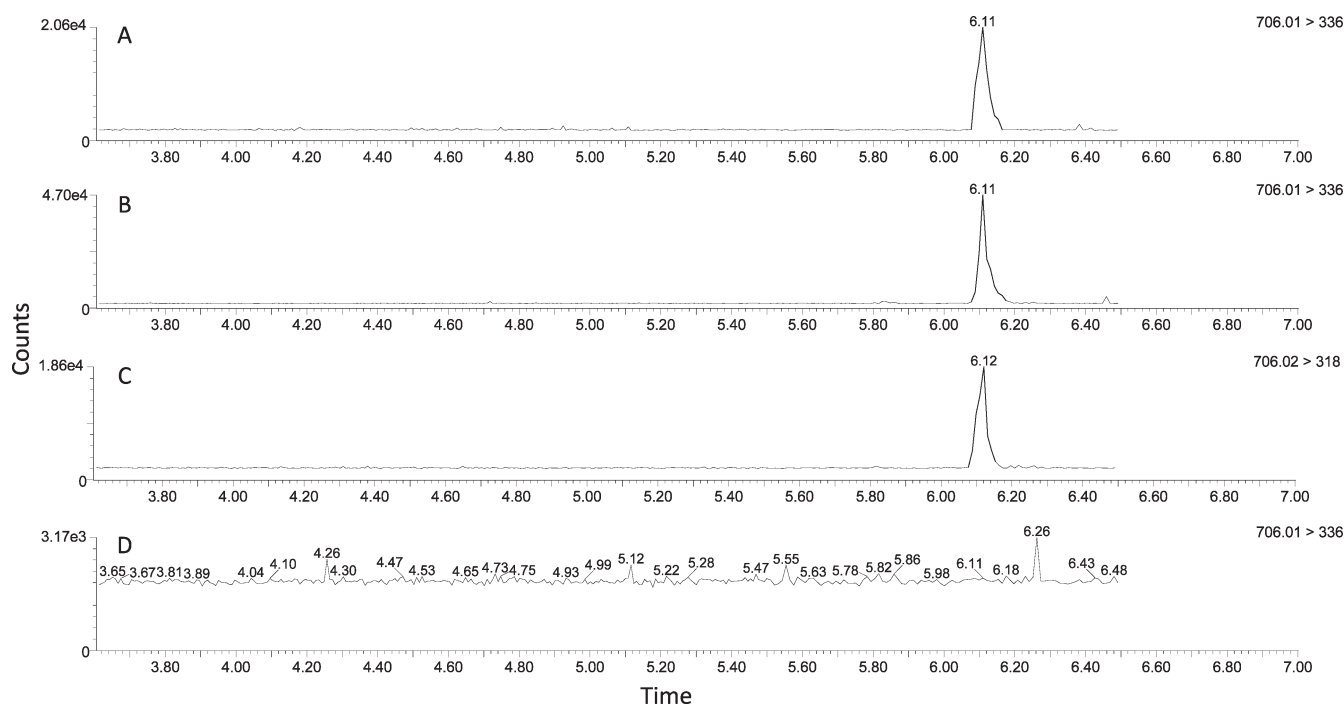


Figure 2. Results from the LC–MS/MS analysis. Transitions of fumonisin B₂ in spiked wine and a wine sample. (A) Quantifier of fumonisin B₂ from spiked wine (45 $\mu\text{g/L}$). (B) Quantifier of fumonisin B₂ from a wine sample (25 $\mu\text{g/L}$). (C) Qualifier of fumonisin B₂ from a wine sample (25 $\mu\text{g/L}$). (D) Quantifier of fumonisin B₂ from a blank sample (0 $\mu\text{g/L}$).

the ¹³C fumonisin B₂, the recovery seemed to be better than 90%. However, clearly, it should not be necessary to use immunoaffinity purification in combination with LC–MS/MS because of the selectivity of the latter technique. The extra purification of 24 samples included 5 control samples (1 blank and 4 spiked with 2–45 $\mu\text{g/L}$). Re-analysis of the 18 initial positive samples did not change the number of positive samples, further confirming that the findings were true positive results. Of the 18 positive samples, 16 were red wine, 1 was white wine, and 1 was port wine (**Table 1**). The positive samples contained 1–25 $\mu\text{g/L}$ fumonisin B₂ (average, 5.2 $\mu\text{g/L}$; median, 2.6 $\mu\text{g/L}$). Selected tested samples are shown in **Figure 2**. On the basis of the relative low amounts of samples, it is indicated that the fumonisin contamination could pose a larger problem in red wine (28% positive) compared to white wine (7% positive). This is similar to the ochratoxin A contamination, which is found to be at a higher concentration and coincidence in red wine compared to white wine (18, 19). The findings of fumonisin B₂ with such a high frequency show that *A. niger* is apparently commonly growing on grapes in the fields. However, the frequency of heavily infected berries was low

because the detected levels are 100–400 lower than those found in highly infected berries (2–8 mg/L) (10, 11). Levels of ochratoxin A in wine have been found to be at 0.002–7.63 $\mu\text{g/L}$ (19, 20), which are lower than the found fumonisin concentrations. There were no statistically significant differences between the fumonisin concentration and countries and years, because of the relatively few samples investigated here.

The maximum permitted level of ochratoxin A in wine is set at 2.0 $\mu\text{g/L}$ by the European Community (EC); this is ca. 2–3 times lower than the average maximum permitted level of ochratoxin A in food (21, 22). Thus, a crude comparison to fumonisin, one could expect a 2–3-fold lower level than the 1–2 mg/kg set by the EC (14) and U.S. FDA (13), respectively. Thus, the maximum level detected (25 $\mu\text{g/L}$) was 40–80 times lower than an anticipated regulatory limit set by the EC and U.S. FDA.

However, even though the detected fumonisin concentrations are below such a limit, it is necessary to perform a larger survey to establish if wines with even higher concentrations are produced, because climatic and production conditions vary. Exposure assessment needs to be performed, especially concerning groups

with a high intake of wine alone or combined with other fumonisin-contaminated products.

In conclusion, this is the first report on the detection of fumonisin B₂ in wine, showing that almost a quarter of the samples were positive. This clearly points toward a much larger survey of how widespread fumonisins are in wine, as well as a determination of the fate of fumonisins in the winemaking process.

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Paper 4

"Occurrence of fumonisins B₂ and B₄ in retail raisins"

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Occurrence of Fumonisin B₂ and B₄ in Retail Raisins

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Concerns that raisins may be contaminated by fumonisins stem from the persistent occurrence of *Aspergillus niger* spores on raisins and the recent discovery of fumonisin production by *A. niger* on grapes, which leads to the widespread occurrence of fumonisin B₂ in wine. This study presents an LC-MS/MS survey of fumonisins in retail raisins. In 10 of 21 brands collected in Denmark, Germany, and The Netherlands, fumonisins B₂ and B₄ were detected at levels up to 13 and 1.3 µg/kg, respectively. Only fumonisin B₂ has been detected in wine, so the presence of fumonisin B₄ in raisins suggests that the fumonisins are produced mainly during the drying process concomitant with the decreasing water activity. Analysis of multiple packages from one manufacturer showed a 3-fold package-to-package variation, suggesting that a few raisins per package are contaminated.

KEYWORDS: Fumonisin B₂ and B₄; wine; grapes; cation exchange; SPE; solid-phase extraction

INTRODUCTION

Black aspergilli are distributed globally and occur on a great variety of substrates. They are also widely used in the food and biotechnology industries for the production of organic acids, enzymes, and other products (1).

Black aspergilli are commonly found on the surface of healthy grapes during all growth stages (2). *Aspergillus* species are present on grape clusters early in the season, and their frequency increases in the later growth stages of the grapes (3). Of the various *Aspergillus* species, *Aspergillus niger* is by far the most commonly found on grapes and is shown in one study to occur on > 80% of samples (4). Although *A. niger* is the predominant species, *Aspergillus carbonarius* is the most problematic because it consistently produces high amounts of ochratoxin A, whereas only 8–40% of *A. niger* strains produce low amounts of this toxin (5,6). This is the main mycotoxin-related health concern in grape-derived products. Nonetheless, *A. niger* production of fumonisins B₂, B₄, and B₆ (7–9) (Figure 1) and the high frequency with which fumonisin B₂ occurs in wine (10, 11) raise concerns about the possible widespread presence of fumonisins in raisins. These concerns are further fuelled by the production of fumonisins B₂ and B₄ on artificially infected grapes and raisins to levels as high as 8 mg/kg (12, 13), as well as a study that found *A. niger* spores on all raisins investigated (13). A recent analysis of seven raisin samples from around the world claimed fumonisin levels as high as 35 mg/kg, mainly for fumonisin B₂, but also detected fumonisins B₁, B₃, and B₆, as well as a number of tentatively identified fumonisin B₁ and B₂ OH-positional isomers (14). The finding of fumonisins B₁ and B₃ is unusual, because the *FUM2* gene encoding the enzyme for a C-10 hydroxylation that is required for the production of these isomers has not been detected in any of

the three available full-genome sequences for *A. niger* strains (15, 16). Furthermore, no other studies on fumonisin production in *A. niger* have detected fumonisin B₁ (7–10, 12, 13, 16–18).

Structurally, fumonisins are similar to sphingolipids and have been shown to inhibit sphingolipid biosynthesis via the ceramide synthase pathway (19). The consumption of contaminated maize-based foods is associated with a high incidence of esophageal and liver cancers, along with neural tube defects (20). Furthermore, outbreaks of leukoencephalomalacia in horses and pulmonary edema and hydrothorax in pigs have been linked to consumption of fumonisin-contaminated feed (20, 21). The regulatory limit for fumonisins in maize is 2–4 µg/g total fumonisins (22) in the United States and 0.2–2 µg/g in the European Union (23). Thus, the findings of 1–36 mg/kg in selected highly infected single raisins and artificially contaminated raisins clearly exceeds regulatory limits (13, 14) and are therefore of public health concern. However, to our knowledge, no studies have reported average fumonisin levels in commercial raisins.

Raisin production methods vary in different parts of the world, but mainly involve sun-drying, shade-drying, or mechanical drying (24). Production of dark-colored raisins in the United States, Chile, and South Africa is by direct sun-drying (24), which usually takes between 2 and 4 weeks to reach the commercial limit of > 85% dry matter (24, 25). Faster drying may be obtained by the use of dipping solutions, commonly used in Mediterranean regions, where grapes are pretreated with solutions of 5–6% K₂CO₃ and olive oil to remove the waxy outer layer prior to sun-drying (26). This practice is generally used to produce the more light-colored sultana raisin variant (24–26). Golden variants are generally mechanically dried after pretreatment with SO₂ to retard nonenzymatic and enzymatic browning (24). Damaged grapes will likely become contaminated, especially by black aspergilli, during the drying period, because these fungi are present on both the grapes and the finished raisins (13). To keep

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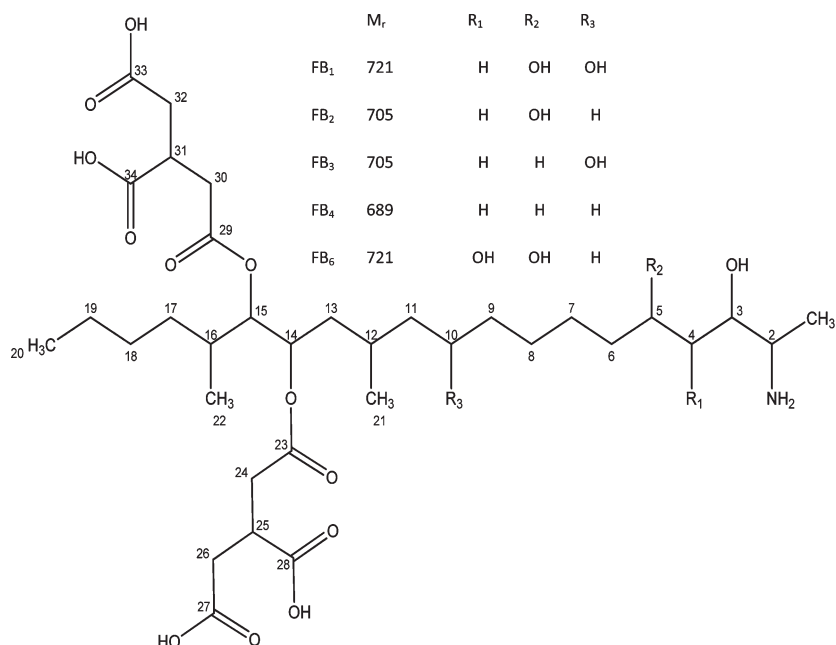


Figure 1. Structure of fumonisins B₁–B₄ and B₆.

ochratoxin A levels low, careful removal of infected raisins is enforced.

In this study, we investigated the presence of fumonisins in commercial raisins using cation-exchange purification with subsequent LC-MS/MS determination. We also investigated various extraction techniques to accommodate the high amounts of interfering material in raisins.

MATERIALS AND METHODS

All solvents were of HPLC grade, and other chemicals were of analytical grade. Reagents were from Sigma-Aldrich (St. Louis, MO). Water was purified on a Milli-Q system (Millipore, Bedford, MA). The fumonisin standard was a mixed certified standard containing fumonisins B₁ at 50.2 mg/L and B₂ at 51.0 mg/L (Biopure, Tulln, Austria). Fumonisin B₃ was from Biopure, and fumonisins B₄ and B₆ were from previous studies (8).

Raisin Samples. Raisins of 21 different brands were purchased from supermarkets and specialty shops in Denmark, Germany, and The Netherlands. The raisins were from the United States/California ($n = 9$), Greece ($n = 1$), South Africa ($n = 2$), Chile ($n = 5$), Turkey ($n = 3$), and China ($n = 1$). The 21 different brands represented a total of six different grape varieties including Thompson seedless, Jumbo Regal seedless, Flame, Sultana, Zante Currant, and Golden seedless.

Optimization of Sample Preparation. Raisins were homogenized with water (1:1 w/w) in a domestic kitchen blender. The mash was subsequently spiked with fumonisins and extracted. For method development, mixtures of water/methanol, water/ethanol, and water/acetonitrile, in 10% stepwise concentration increases from 30 to 100% organic solvent, were tested and the extracts purified using several formats (30 or 60 mg in 1 or 3 mL) of Strata X-C columns (Phenomenex, Torrance, CA). The effects of formic acid addition during extraction and different extraction volumes (3–20 times the amount of mash) were tested, along with different concentrations of NH₄OH for elution from the cation-exchange columns.

Sample Preparation. Raisins (≥ 200 g) were homogenized as described above, and 34 g of mash was transferred and split into two 50 mL Falcon tubes. The homogenates in each tube were extracted with 40 mL of acetonitrile/water/formic acid (86:12:2 v/v/v) under continuous shaking for 2 h. Samples were then centrifuged at 8000g for 5 min, and 35 mL of the upper phase of the resulting two-phase system was transferred to a clean 50 mL conical tube. Samples were evaporated to dryness under a N₂ flow, and each was redissolved in 10 mL of 1:1 v/v methanol/water, passed through a 0.45 μ m PTFE syringe filter, and pooled. Sample purification was carried out using 60 mg/3 mL Strata X-C columns that had been

preconditioned with 1 mL of methanol and 1 mL of water prior to sample load. To further acidify the column, 1 mL of water containing 2% formic acid was added, with a subsequent wash with 1 mL of methanol. Fumonisins were eluted with 1 mL of methanol/2% NH₄OH (1:4 v/v) into an autosampler vial and used directly for analysis.

LC-MS/MS conditions. LC-MS/MS analysis was performed as previously described (11, 13) on a Quattro Ultima triple mass spectrometer (Micromass, Manchester, U.K.) using positive electrospray ionization and operated in multiple-reaction mode (MRM or SRM) in the following transitions: fumonisins B₂ and B₃, quantifier m/z 706 \rightarrow 336, cone, 50 V; collision, 40 eV; dwell time, 50 ms; qualifier m/z 706 \rightarrow 318; cone 50 V; collision, 25 eV; dwell time, 100 ms; fumonisin B₄, quantifier m/z 690 \rightarrow 320; cone, 50 V; collision, 35 eV; dwell time, 50 ms; qualifier m/z 690 \rightarrow 338; cone, 50 V; collision, 30 eV; dwell time, 100 ms; fumonisins B₁ and B₆, quantifier m/z 722 \rightarrow 334; cone, 50 V; collision, 40 eV; dwell time, 50 ms; qualifier m/z 722 \rightarrow 528; cone, 50 V; collision, 25 eV; dwell time, 100 ms.

Calibration. Quantification was by external standard quantification using linear regression, with comparison to spiked raisin samples (raisins/water 1:1 w/w) incubated for 2 h prior to extraction. Samples were spiked with a certified mixture of fumonisins B₁ and B₂ to 0 (blank), 1, 5, 15, 50, or 100 μ g/kg dry raisins. Preliminary results had indicated that the samples had low fumonisin levels that were close to the limit of quantitation (LOQ), so each homogenized sample was analyzed in triplicate. Because available amounts of fumonisins B₄ were low, its response factor relative to fumonisin B₂ was determined on the basis of pure standards.

Validation. Fumonisin B₂ determination was validated by spiking portions of 34 g of raisin mash (raisin/water 1:1 w/w) with a mixture of fumonisins B₁ and B₂ to concentrations of 0 (blank), 1, 5, 15, 50, or 100 μ g/kg dry raisins. Experiments were performed on three different days, with all levels assayed in triplicate each day. Apparent recovery was calculated as the ratio of the slope of the spiked raisins to a standard dilution series in methanol/water (20:80 v/v) diluted to 0, 1, 5, 15, 50, and 100 μ g/kg. Fumonisins B₆ and B₃ were occasionally coanalyzed in sequence to ensure sufficient separation.

RESULTS AND DISCUSSION

Sample Purification. Fumonisins have traditionally been purified using anion-exchange or immunoaffinity columns; however, in previous studies we observed that raisin extracts fouled immunoaffinity columns, probably due to polysaccharides and brown pigments (13), resulting in brownish viscous methanol eluates. Poor recovery has been observed when using anion-exchange columns for grape extracts and wine, most likely

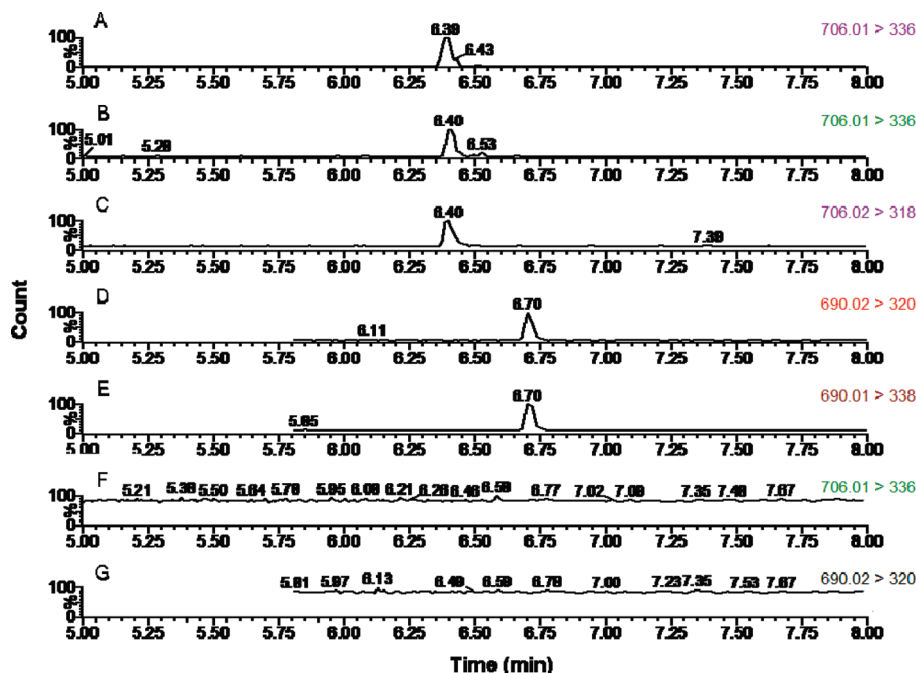


Figure 2. LC-MS/MS analysis; fumonisin B₂ and B₄ transitions in a fumonisin B₂-spiked sample and a blank raisin sample: (A) quantifier of fumonisin B₂ in a spiked raisin sample (5 μ g/kg); (B) quantifier and (C) qualifier of fumonisin B₂ in a raisin sample (4.6 μ g/kg); (D) quantifier and (E) qualifier of fumonisin B₄ in a raisin sample (1.2 μ g/kg); (F) quantifier of fumonisin B₂ in a noncontaminated raisin sample; (G) quantifier of fumonisin B₄ in a noncontaminated sample.

because of the high content of organic acids, which salt-out fumonisins (11, 13). To circumvent this problem, we used polymeric mixed-mode cation reversed phase columns, which target the amine of the fumonisins and have been proven effective for purifying fumonisins from wine (11).

During method development, we tested ethanol/water and methanol/water mixtures for extraction, because they did not result in a two-phase system as acetonitrile/water did, as also observed for aflatoxin extraction in various dry sample types (27). However, extensive amounts of pigments and polysaccharides (as judged from the extract viscosity) subsequently interfered with solid-phase extraction column function. Because this did not occur using acetonitrile extracts, we selected this QuEChERS-like method (28) in which the acetonitrile/water phase separates because of sugars from the raisins and performed further optimizations on this. Not surprisingly, the extracts needed to be acidified to keep the fumonisins in the acetonitrile phase. We also found that the volume of acetonitrile increased with decreasing temperature (observed when extracted stored in the refrigerator), so the samples were processed at the same temperature as the spiked samples used for calibration. We used two 50 mL conical tubes instead of a single 100 mL tube for extraction and extract evaporation to fit our centrifuge and N₂ evaporation equipment.

Validation. The LOQ was determined to be 1 μ g/kg for fumonisin B₂ at a relative standard deviation of 35% ($n = 9$), which is highly satisfactory for this low level in this type of matrix. Trueness was 89, 101, and 97% for the three days, with average deviations of 22, 8, and 9%. The limit of detection (LOD, $s/n = 5$) was at least 0.3 μ g/kg for fumonisin B₂ and 0.3 and 0.1 μ g/kg for fumonisin B₄. Apparent recovery was $87 \pm 15\%$.

Screening of Raisin Samples. Our analysis demonstrated that 48% of the 21 tested raisin brands contained fumonisin B₂ and 43% fumonisin B₄. The detected amounts of fumonisins B₂ and B₄ in the positive samples ranged from the LOD (0.3 and 0.1 μ g/kg) to 13 and 1.3 μ g/kg, respectively. LC-MS/MS MRM chromatograms from selected spiked and nonspiked samples are shown in Figure 2.

The detection of fumonisin B₄ from retail raisins matches previous studies on artificially infected grapes and raisins (13); however, fumonisin B₄ was not identified in studies of fumonisin content in wine and grape must (10–12). This might suggest a different growth physiology and metabolism of *A. niger* on the drying grapes, possibly because of a fast fungal growth rate that results in some fumonisin B₄ not being transformed into B₂ before the water activity decreases below *A. niger* growth limits (~ 0.85) (4). This postharvest growth along with the physical removal of water from the vine fruit and the concomitant concentration of the fumonisins may explain why fumonisin B₄ has not been detected in previous wine studies (10, 11). The presence of fumonisin B₄ in raisins could also explain our higher frequency of positive samples because, in contrast to wine production, both the postharvest period and the drying are sources of fumonisin production and upconcentration.

The frequency of fumonisin contamination in wine is approximately 18–23% (10, 11), which is half the frequency observed in the raisins in this study, fitting well with the higher frequency being caused by postharvest growth of *A. niger*. This correlates well with studies demonstrating a significant increase in fungal load during drying, exceeding the expected increase due to the concentration effect of water evaporation (4).

The levels we detected were > 1000-fold lower than in samples analyzed by Varga et al. (14), in which fumonisin B₁ was also detected. The finding of fumonisin B₁ is highly unlikely because bioinformatic analysis of the data from the three fully sequenced *A. niger* strain genomes (15) has not detected a *FUM2* gene encoding the enzyme responsible for C-10 hydroxylation (29). Thus, *A. niger* should not produce fumonisins B₁ and B₃ as also reported here and in all other studies analyzing extracts from *A. niger* strains (7–10, 12, 13, 16–18).

Of the 10 positive samples in this study, 7 were derived from California ($n = 9$), 2 from Chile, and 1 from Turkey (Table 1), demonstrating a higher infection rate in the California raisins. The reasons for this are unknown and could be because of the different climate, preharvest weather conditions, different grape

Table 1. Overview of Tested Raisin Samples and Average Fumonisin Content^a

brand	origin	grape variety	organic	fumonisin B ₂ (μg/kg)	fumonisin B ₄ ^b (μg/kg)
1	California	Thompsons Seedless		4.9 ± 0.4	0.26 ± 0.05
1				4.5 ± 1	0.8 ± 0.4
1				13.0 ± 1	1.3 ± 0.4
1				7.5 ± 0.8	0.9 ± 0.2
2	California	Thompsons Seedless		2.7 ± 0.2	0.29 ± 0.1
3	California	Thompsons Seedless	✓	1.8 ± 0.2	<LOQ ^c
4	California	Thompsons Seedless	✓	2.6 ± 0.4	0.83 ± 0.5
5	California	Thompsons Seedless		3.8 ± 0.6	0.31 ± 0.09
6	California	Thompsons Seedless		4.1 ± 0.6	1.0 ± 0.1
7	California	Thompsons Seedless	✓	1.3 ± 0.6	0.25 ± 0.2
8	California	Thompsons Seedless		ND ^d	ND
9	California	Thompsons Seedless	✓	ND	ND
10	Turkey	Sultana	✓	ND	ND
11	Turkey	Sultana	✓	ND	ND
12	Turkey	Sultana		4.8 ± 2	0.75 ± 0.1
13	Greece	Zante currants		ND	ND
14	South Africa	Golden Seedless		ND	ND
15	South Africa	Jumbo Regal Seedless	✓	ND	ND
16	China	unknown		ND	ND
17	Chile	Flame		ND	ND
18	Chile	Flame		ND	ND
19	Chile	Flame		ND	ND
20	Chile	Golden		<LOQ	ND
21	Chile	Golden		1.4 ± 0.6	<LOQ

^a Fumonisin concentration is the average of three determinations from the same package, ± one standard deviation. ^b Fumonisin B₄ was calculated assuming a 4-fold better response factor than for fumonisin B₂ and is shown from one spiking experiment. Fumonisin B₁ and B₆ were not detected in any of the packages. ^c LOQ, limit of quantification: for fumonisin B₂, 1 μg/kg; for fumonisin B₄, estimated as 0.25 μg/kg. ^d ND, not detected. The limit of detection for fumonisin B₂ was 0.3 μg/kg, and that for fumonisin B₄ was 0.1 μg/kg.

cultivars, or different drying and sorting techniques (4). No connection was observed between fumonisin content and type of cultivation strategy (organic/conventional).

Large package variations were observed with up to 3-fold differences between four packages of the same brand, indicating a nonhomogeneous infection level, which may be due to contaminated raisins. Even though the detected fumonisin levels were 50–100 times below the regulatory limits of similar commodities such as maize, a larger survey is needed, both because of the low number of retail brands tested and because of the high variation among packages. Furthermore, establishing whether raisins may have even higher concentrations under different climatic conditions and drying procedures is important.

We presume that the low levels that we report here are also because of the efficient removal of damaged grapes that was initiated after problems with ochratoxin A in grapes and derived products were reported in the late 1990s (30–32), leading to very strict European Community regulations (472/2002) (33), including a maximum allowance of 10 μg/kg ochratoxin A in dried vine fruits (34).

In conclusion, this first survey on fumonisin determination in retail raisins demonstrated that almost 50% of tested brands contained low levels of fumonisins (1–14 μg/kg), which is far from the maximum allowable levels in regulated foods such as maize. Nonetheless, our finding of a high frequency of fumonisin occurrence clearly calls for larger surveys that can establish whether certain regions or weather conditions lead to fumonisin concentrations above set regulatory limits, as well as investigations of the influence of the drying process, especially on the production of fumonisin B₄.

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Paper 5

" *Aspergillus acidus* from Puerh tea and black tea does not produce ochratoxin A and fumonisin B₂"

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Aspergillus acidus from Puerh tea and black tea does not produce ochratoxin A and fumonisin B₂

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ABSTRACT

Puerh tea is a unique Chinese fermented tea. Unlike other teas it is stored for a long period of time. *Aspergillus niger* is claimed to be the dominant microorganism in the Puerh tea manufacturing process and also to be common on tea in general. *A. niger sensu stricto* is known to produce the mycotoxins ochratoxin A, fumonisins B₂ and B₄. With this in mind, we performed a preliminary study to determine if production of these mycotoxins by black *Aspergilli* isolated from Puerh and black tea can occur. An examination of 47 isolates from Puerh tea and black tea showed that none of these was *A. niger*. A part of the calmodulin gene in 17 isolates were sequenced, and these 17 isolates were all identified as *Aspergillus acidus* (= *A. foetidus* var. *acidus*). The rest of the 47 isolates were also identified as *A. acidus* from their metabolite profile. Neither production of ochratoxin A nor fumonisins B₂ and B₄ by any of the 47 isolates were observed.

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1. Introduction

Puerh tea is a black tea, which is produced mainly in the Chinese Yunnan district and consumed by a large part of the Chinese population (Sano et al., 1986). The Puerh tea is different from other teas, in the sense that it is fermented as a normal black tea, but is matured for a period of time (Sano et al., 1986). It is said that the longer the maturation, the better taste and quality of the Puerh tea (Sano et al., 1986). According to the literature *Aspergillus niger* is associated with black tea and is the dominating microorganism during the Puerh tea production (Abe et al., 2008; Elshafie et al., 1999; Mo et al., 2005; Sano et al., 1986; Xu et al., 2005). *A. niger* and other members of section *Nigri* are difficult to classify and identify because they are morphologically very similar (Samson et al., 2007). Physiology, secondary metabolites and DNA sequencing can help to differentiate between the species (Samson et al., 2004; Samson et al., 2007).

Recently a putative fumonisin gene cluster was discovered in two genomes of *A. niger*, strains ATCC 1015 and CBS 513.88 (Baker 2006; Pel et al., 2007). This was followed by the discovery of the ability to produce fumonisins B₂ (FB₂) and B₄ (FB₄) by *A. niger* (Frisvad et al., 2007; Noonim et al., 2009). Besides this production *A. niger* is also a well known ochratoxin A (OTA) producer (Abarca et al., 1994). Fumonisin B₁ has previously been isolated from black tea (Martins et al., 2001), and since Puerh tea is presumably fermented with *A. niger*, there is a possibility that fumonisins and potentially OTA from *A. niger* can be found in Puerh tea.

We therefore investigated the ability of 47 black *Aspergilli*, isolated from black tea and Puerh tea, to produce OTA as well as FB₂ and FB₄.

2. Materials and methods

Unless otherwise stated all solvents were HPLC grade, chemicals were analytical grade and water was purified on a Milli-Q system (Millipore, Bedford, MA). Media were prepared in 9 cm Petri dishes, each with 20 ml substrate, and strains were inoculated by the three point method and dishes incubated in micro perforated plastic bags for 7 days in darkness. All samples were as a minimum made in biological duplicates on two separate plates.

2.1. Samples

Ten different tea samples were purchased from the internet (www.tea.dk) or in supermarkets/specialty shops in Denmark or The Netherlands. All teas were end products, 5 were Puerh teas (Brand A–E) and 5 were regular black teas (Brand 1–5). The Puerh teas were all from the Yunnan Region, China. The black teas were from Anhui region (China), Singtom (India) or Sri Lanka.

2.2. Screening

One gram portions of each tea were taken at random from well mixed portions (min. 50 g). Afterwards, each portion was plated onto five Petri dishes, containing dichloran 18% glycerol agar (DG18) (Hocking and Pitt, 1980). The plates were incubated for seven days at 25 °C. After

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incubation, up to five black *Aspergillus* cultures from each tea were isolated and inoculated on Czapek yeast autolysate agar with 5% salt (CYAS) (Frisvad and Samson, 2004), malt extract agar (Pitt, 1979), yeast extract sucrose agar (YES) (Frisvad and Samson, 2004) and two Czapek yeast autolysate agar (CYA) (Frisvad and Samson, 2004). All plates were incubated for seven days at 25 °C. In addition a CYA plate incubated at 37 °C was used for each isolate. Morphological and physiological studies were performed using all five plates.

2.3. Analysis of secondary metabolites

Three plugs ($D=6$ mm) from the CYA (25 °C) and three plugs ($D=6$ mm) from the YES plate were transferred to a 2 ml dram vial and 800 μ l of methanol:dichloromethane:ethyl acetate (1:2:3; vol:vol:vol) with 1% formic acid was added. The plugs were extracted ultrasonically for one hour. The extract was transferred to a new vial and the organic phase was evaporated in a rotational vacuum concentrator. The remains were redissolved by ultrasonication in 500 μ l methanol for twenty minutes. The samples were filtrated through a FTPE 0.45 μ m filter and then analyzed by HPLC-DAD (Smedsgaard et al., 1997). The potential FB₂

and FB₄ production were investigated by the extraction method described by Frisvad et al. in 2007 with small modifications. Extraction of two separate sets of six plugs from the CYAS and YES plate were performed. The plugs were extracted ultrasonically for one hour with 800 μ l methanol:water (3:1). The extract was filtrated through a FTPE 0.45 μ m filter and used directly for LC-MS.

2.4. HPLC conditions

Extracts were analyzed on a Agilent 1100 HPLC system equipped with a diode array detector. Scan: 190–600 nm with a bandwidth of 4 nm approx. two times per second. The separation was performed on a 100 \times 2 mm, 3 μ m particle size, i.d. Luna C-18 (II) column fitted with a 4 \times 2 mm, 3 μ m particle size, C18 security guard column. Using a water:acetonitrile, the gradient started at 15% acetonitrile and was increased to 100% acetonitrile in twenty minutes and kept there for five minutes, before it returned to starting conditions. The flow rate was 0.3 ml/min. Both eluents contained 0.005% (v/v) trifluoroacetic acid (modified from Smedsgaard et al., 1997). All chemicals were analytical grade. 3 μ l extract was injected for all samples.

Table 1

Known metabolites produced by 47 black *Aspergilli* isolated from tea.

IBT no.	Type of tea (Brand)	Asperazine	Funalone	Pyranonigin A	Naphto- γ -pyrones	Antafumicin	FB ₂	FB ₄	OTA	Identified as
24798	Black tea (1)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24799	Black tea (1)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24800	Black tea (1)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24801	Black tea (2)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24802	Black tea (2)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24803	Black tea (2)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24821	Black tea (3)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24822	Black tea (3)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24825	Black tea (3)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24826	Black tea (3)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
24830	Black tea (3)	X	X	–	X	X	–	–	–	<i>A. acidus</i>
24831	Black tea (3)	X	X	X	X	X	–	–	–	<i>A. acidus</i>
29563	Black tea (4)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29564	Black tea (4)	X	X	–	X	–	–	–	–	<i>A. acidus</i>
29565	Black tea (4)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29566	Black tea (4)	X	X	–	X	–	–	–	–	<i>A. acidus</i>
29567	Black tea (4)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29568	Puerh tea (A)	X	X	–	X	–	–	–	–	<i>A. acidus</i>
29569	Puerh tea (A)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29570	Puerh tea (A)	X	X	–	X	–	–	–	–	<i>A. acidus</i> ^a
29571	Puerh tea (A)	X	X	–	X	–	–	–	–	<i>A. acidus</i> ^a
29572	Puerh tea (B)	X	X	X	–	–	–	–	–	<i>A. acidus</i>
29573	Puerh tea (B)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29574	Puerh tea (B)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29575	Puerh tea (B)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29576	Puerh tea (B)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29577	Puerh tea (C)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29578	Puerh tea (C)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29579	Puerh tea (C)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29580	Puerh tea (C)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29581	Puerh tea (C)	X	X	–	X	–	–	–	–	<i>A. acidus</i>
29582	Puerh tea (D)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29583	Puerh tea (D)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29584	Puerh tea (D)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29585	Puerh tea (D)	X	X	X	–	–	–	–	–	<i>A. acidus</i> ^a
29586	Puerh tea (D)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29587	Black tea (5)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29588	Black tea (5)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29589	Black tea (5)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29590	Black tea (5)	X	X	–	X	–	–	–	–	<i>A. acidus</i>
29591	Black tea (5)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29592	Puerh tea (E)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29593	Puerh tea (E)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29594	Puerh tea (E)	X	X	X	X	–	–	–	–	<i>A. acidus</i>
29595	Puerh tea (E)	X	X	–	–	–	–	–	–	<i>A. acidus</i>
29596	Puerh tea (E)	X	X	X	X	–	–	–	–	<i>A. acidus</i> ^a
29597	Puerh tea (A)	X	X	X	X	–	–	–	–	<i>A. acidus</i>

^a Identified from molecular data.

2.5. LC-MS conditions

The LC-MS analysis was performed on a LC/MSD VL single quadrupole (Agilent, Santa Clara, California, USA). The separation was done on a 50 × 2 mm, i.d. 3 µm particle size, Luna C-18 (II) column (Phenomenex, Torrance, California), which was heated to 40 °C, fitted with a security guard column, using a water:acetonitrile (both containing 20 mM formic acid) gradient at a flow rate of 0.3 ml/min. The gradient started at 30% acetonitrile, and increased to 60% acetonitrile over 5 min. During further 1 min it was increased to 100% acetonitrile and maintained for 2 min before the gradient was returned to starting conditions in 1 min and kept there for 5 min. A 3 µl extract was injected for all samples. The mass spectrometer (MS) was operated in positive electrospray ionization mode and was automatically calibrated on the instrument ESI tuning mix (Agilent). The MS was used in selected ion monitoring (SIM) mode for the following $[M+H]^+$: FB₂ (m/z 706), FB₄ (m/z 690). The capillary voltage was held at 3000 V, the fragmentor voltage was at 70 V and the nebuliser pressure was at 2.5 bar. The drying gas flow was 12 l/min with a temperature of 350 °C. Selected extracts were verified by LC-DAD-HRMS (Frisvad et al., 2007).

2.6. Isolation and analysis of nucleic acids

The cultures used for the molecular studies were grown on malt peptone (MP) broth using 10% (v/v) of malt extract (Oxoid) and 0.1% (w/v) bacto peptone (Difco), 2 ml of medium in 15 ml tubes. The cultures were incubated at 25 °C for 7 days. DNA was extracted from the cells using the Masterpure™ yeast DNA purification kit (Epicentre Biotechnology, Madison, Wisconsin) according to the instructions of the manufacturer. Amplifications of the partial calmodulin gene were set up as described previously (Hong et al., 2006). All the sequencing reactions were purified by gel filtration through Sephadex G-50 (Amersham Pharmacia Biotech, Piscataway, NJ) equilibrated in double-distilled water and analyzed on the ABI PRISM 310 Genetic Analyzer (Applied Biosystems) using the Big Dye Terminator Cycle Sequencing Ready Reaction Kit for both strands, and the sequences were aligned with the MT Navigator software (Applied Biosystems).

3. Results and discussion

In total 47 black *Aspergilli* were isolated from the tea samples. The metabolite production of the 47 black *Aspergilli* was analyzed (Table 1). None of the 47 isolates was able to produce FB₂, FB₄ nor OTA. All of the 47 isolates produced asperazine and funalenone. In addition, production of pyranonigin A was observed in 38 of the isolates, naphto-γ-pyrone in 44 isolates and antafumicins in 12 isolates. According to the literature, *A. niger sensu stricto* is able to produce both funalenone, naphto-γ-pyrone and pyranonigin A, but lacks the ability to produce antafumicin and asperazine and this species was hereby excluded as being the species from the tea (Samson et al., 2007). Since *A. foetidus* has recently been synonymized with *A. niger* (Peterson 2008) only three members within section *Nigri* are able to produce asperazine. These are *Aspergillus acidus* (= *A. foetidus* var. *acidus*; (Kozakiewicz 1989)), *A. tubingensis* and *A. vadensis* (Samson et al., 2007). The latter, *A. vadensis*, is not able to produce either antafumicin, funalenone or pyranonigin A (Samson et al., 2007). However, 12 of the isolates were able to produce antafumicins, which are only produced by *A. acidus* (Samson et al., 2007). The last 35 isolates were hereby provisionally identified as *A. acidus* or *A. tubingensis*. To further help the identification process we examined sequences of part of the calmodulin gene in 17 isolates, randomly chosen from these 35 isolates to prove their species assignment. The partial calmodulin gene sequences of all the examined isolates were identical to that of the type strain of *A. acidus*, isolate CBS 564.65^T (= ITEM 4507 = ATCC 16874 = IMI 104688; Genbank accession number AM19749). All 35 isolates were hereby identified as *A. acidus*.

Even though we have a limited amount of samples the results from this preliminary study point to the fact that *A. acidus* is the dominant fungus in Puerh tea and black tea, in contrast to previous studies which describe *A. niger* as the dominant fungus in Puerh tea fermentation and in black tea (Abe et al., 2008; Bouakline et al., 2000; Mo et al., 2005; Sano et al., 1986; Xu et al., 2005). In these studies a broader concept of *A. niger* was used, which would include what is circumscribed as *A. acidus* today. This leads us to conclude that the reported black *Aspergilli* from Puerh tea and black tea probably should be re-identified as *A. acidus*. In our screening there was no indication of *A. niger sensu stricto* in any of the tea analyzed in this study. Since only *A. acidus* is found and this does not produce FB₂, FB₄ nor OTA, it is very unlikely that the production of these mycotoxins by *A. niger sensu stricto* can occur in tea. This correlates with the observation that among the fumonisins, only FB₁ from *Fusarium* spp. has been isolated from tea, and *A. niger* is not able to produce FB₁ (Frisvad et al., 2007; Martins et al., 2001).

To support this preliminary study, further investigation of more samples, as well as additional samples from different regions and stages in the tea processing should be performed. However, it is predicted that no or a very low amount of *A. niger* will be found in Puerh tea or other black teas, making any presence of FB₂, FB₄ and OTA from *A. niger* in this commodity less likely.

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Paper 6

”Studies on fumonisins in peanuts and peanut butter”

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Paper to be submitted to Journal of Agricultural and Food Chemistry

1 **Studies on fumonisins in peanuts and peanut butter**

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8

9 **Abstract**

10 Fumonisin are important carcinogenic mycotoxins, mainly associated to maize and other crops
11 infected with *Fusarium* species. Recently *Aspergillus niger* contamination have been shown to be
12 the source of fumonisin contamination in grapes, raisins and wine, and we speculate if peanut
13 derived products were also contaminated by fumonisins as *A. niger* is a very frequent contaminant
14 of peanuts. Here we investigate the production of fumonisins on both peanuts artificially
15 contaminated by *A. niger*, as well as retail peanuts and peanut butter. Fumonisin were quantified
16 by isotope dilution LC-MS/MS after an-ion exchange purification. Fumonisin were detected in all
17 artificially infected peanuts at levels up to 10 mg/kg whereas no fumonisins were detected in 15
18 peanut and 9 peanut butter retail products (limit of detection 0.004 mg/kg). We speculate that the
19 absence of fumonisins in the retail products is a results of the severe damage induced to the nuts
20 by *A. niger* which facilitates easily separation of infected and non-infected nuts.

21

22 Introduction

23 Mycotoxins produced by some fungal species, especially *Aspergillus*, *Penicillium* and *Fusarium*¹
24 are naturally occurring in many food and feed types. The most toxic and commonly occurring ones
25 (aflatoxins, ochratoxin A, fumonisins, trichothecenes, and zearalenone) are subsequently
26 regulated and routinely monitored.

27 The fumonisins (**Figure 1**), are mainly associated with *Fusarium* infected maize but however newer
28 discoveries have shown that also *Aspergillus niger* and three *Tolypocladium* species produce these
29 compounds²⁻⁵. While *Fusarium* species have a wide fumonisin profile showing FB₁-FB₅ as well as
30 the A-, C- and P- groups, *A. niger* is only known to produce FB₂, FB₄ and FB₆ and *Tolypocladium*
31 only FB₂ and FB₄^{2,3,5,6}.

32 Since fumonisins are regularly found in toxic levels in especially maize products they are regulated
33 in to levels of 0.2-4 mg/kg in US and EEC depending on the maize type and intended use^{7,8}.

34 *A. niger* is among the commonest fungi isolated from nuts, especially peanuts, and has also been
35 reported in pecans, pistachios, hazelnuts, walnuts, coconut copra, cereals and oilseeds⁹⁻¹⁴.

36 Infection of peanuts by *Aspergillus*, especially members of section *Flavi* and *Nigri* occurs under
37 both pre-harvest and post-harvest conditions^{15,16}. The main problem organisms concerning
38 peanuts and mycotoxins are *A. flavus* and the related *A. parasiticus*, due to their production of the
39 highly carcinogenic aflatoxins and the fact that they do not visually damage the nuts but only give
40 some discoloration that can be further enhance after roasting¹⁷. Although, *A. niger* is also
41 frequently isolated from peanuts it has until recently not been considered a safety problem as very
42 few strains produce ochratoxin A. But with the discovery that 75-80% of all strains, independent of
43 isolation commodity, can produce fumonisins^{2,3,18-21} peanuts is a subsequently potential source of
44 fumonisin exposure.

45 In a project screening novel fumonisin contaminated food products we decided to: i) investigate if
46 *A. niger* can produce fumonisins while growing on peanuts and ii) develop a LC-MS/MS method for
47 quantification fumonisins in peanuts and peanut butter.

48

49 **Materials and Methods**

50 Unless otherwise stated, all solvents were high-performance liquid chromatography (HPLC)-grade
51 and other chemicals were analytical-grade from Sigma-Aldrich (St. Louis, MO). Water was purified
52 on a Milli-Q system (Millipore, Bedford, MA). The mixed certified standard of U-¹³C₃₄-fumonisin B₂,
53 FB₁ and FB₂ were from Biopure (Tulln, Austria). Fumonisins FB₄ and FB₆ were available from
54 previous studies ⁴.

55

56 **Peanut samples**

57 Peanut samples (n=15) were purchased at local markets from three locations in Thailand (Surat
58 Thani, Chang Mai and Bangkok). Peanut butter samples were purchased from Danish
59 supermarkets and specialty shops. In total 9 brands were tested, they were produced in USA (n=2),
60 the Netherlands (origin unknown) (n=2), Argentina (n=2), Denmark (n=2) (origin: China) and
61 England (n=1) (origin: Indonesia).

62

63 **Inoculation of *A. niger* on peanuts**

64 Shelled peanuts were sterilized in 2% hypochlorite for 2 min, and washed twice in sterile water.
65 The peanuts were placed in a petri dish with water agar. A hole was punched in the shell with a
66 sterile needle (D=1-2 mm). Spore suspension (10 µL) of was placed upon the hole in the peanut
67 shell. The peanuts were then incubated at 25 °C for 7-14 days in darkness. Afterwards a single

68 peanut carefully removed from the shell and extracted. The following four fumonisin producing *A.*
69 *niger* strains, were randomly selected from the IBT collection (Authors address): IBT 18741, 28753,
70 29019 and 29331. Unfortunately no *A. niger* strains originating from peanuts were available, we
71 therefore randomly selected 4 strains from raisins.

72

73 **Sample preparation**

74 One shelled infected/control peanut (cut out of the shell) or 3 g retail peanuts were placed in a
75 cryo tube with 10 steel balls and 3 ml methanol/water, and shaken for 5 minutes by a Mini Bead-
76 beater (Biospec Products Inc, Bartlesville, OK). The peanut butter was considered homogeneous
77 and 3 g peanut butter was extracted directly.

78 The 3 g retail peanuts (or single nuts) were extracted in total with 15 ml methanol/water (3:1 v/v)
79 and 8 ml heptane, by thoroughly shaking for 15 minutes in a 50 ml Falcon tube on a shaking table
80 at 200 RPM. Phases were separated by centrifugation at 9400 rpm for 5 minutes and the liquid
81 middle phase (approx. 12 ml) from the resulting three-phase system was transferred to a new 50
82 ml falcon tube. The extract was further defatted by adding 8 ml heptane, vortexed and centrifuged
83 at 9400 rpm for 5 minutes. The lower phase was transferred to a new falcon tube and excess
84 heptane was removed after additional centrifugation. The pH was measured and adjusted to $6.3 \pm$
85 0.2 with formic acid or ammonium hydroxide. From each sample 9 ml was applied to a 200 mg/3
86 ml Strata SAX column (Phenomenex, Torrance, CA), which was previously conditioned with 1 ml
87 water and 1 ml methanol. The cartridge was afterwards washed with 3 ml of methanol/water (3:1
88 v/v) and 2 ml methanol. Fumonisins were eluted with 2 ml methanol with 1 % formic acid and 20
89 ng U-¹³C₃₄-fumonisin was added to each sample. Solvent was evaporated at 37 °C under a gentle

stream of nitrogen and redissolved in 2x100 µl acetonitrile/water (3:7 v/v) and used directly for analysis.

92

93 **Validation**

94 The fumonisin B₂ extraction were only validated on peanut butter. Validation was performed by
95 spiking portions of 3 g peanut butter to the following concentrations of FB₂: 0 (blank), 0.004,
96 0.008, 0.016, 0.060, 0.25, 1, 4, 16 mg/kg. Three experiments of each validation were performed on
97 3 different days, with all levels in triplicates each day. To determinate the apparent recovery, the
98 ratio of signal of FB₂ for spiked samples and standards was compared without U-¹³C₃₄-FB₂
99 correction.

100

101 **LC-MS/MS conditions**

102 LC-MS/MS analysis was performed as previously described^{20,22}. Briefly described the LC-MS/MS
103 analysis was performed on a Quattro Ultima triple mass spectrometer (Micromass, Manchester,
104 UK) using positive electrospray ionization. Separations was performed on a 50 × 2 mm i.d., 3 µm
105 Gemini C6-phenyl column (Phenomenex, Torrance, CA) as described previously. One thing
106 differentiated from the previously described method, an injection volume of 7 µL was used.
107 Fumonisin B₂, B₄, B₁ and B₆ were monitored using two transitions each^{20,22} while the internal
108 standard, U-¹³C₃₄-FB₂, was monitored by m/z 740 → 358.

109

110 **Results & Discussion**

111 In order to get a robust defatting, several defatting agents were tested, e.g. ethyl acetate,
112 heptane, pentane and toluene. Pentane evaporated too quickly, toluene gave unclear extraction

113 phase and ethyl acetate gave a lower recovery due to emulsion formation, therefore heptane was
114 chosen. At first the defatting and extraction step was divided, but in order speed up the process,
115 the extraction and defatting steps were combined in one step, with no significant drop in recovery.
116 Two different extraction solvents were also tested, MeOH:H₂O (3:1 v/v) and ACN:H₂O (3:1 v/v).
117 The 75% MeOH solution gave similar recovery as the 75% CAN, however ACN was deselected due
118 to sedimentation (data not shown), and MeOH was chosen as the extraction solvent. Robustness
119 testing of the fumonisin extraction at different pH values (5, 7, 9) gave the same recovery (data
120 not shown).

121

122 **Validation**

123 The three calibration curves, one from each day, from spiked samples, had the following R² values:
124 0.9998, 0.9996, and 0.9999. The relative standard deviation (RSD) of the slope (peak area of
125 quantifier/peak area-U-¹³C₃₄ FB₂) from the 3 days varied within 4 and 17% (average 7%). Limit of
126 quantification was 0.008 mg/kg (RSD: 16%) and limit of detection (s/n of 5:1) was 0.004 mg/kg.
127 Absolute recovery of FB₂ was found to be 69 ± 14%.

128

129 **Fungal growth on peanuts – Worst case scenarios**

130 Since *A. niger* is a common contaminant of peanuts^{11,13,16,23,24}, we made worst case scenarios, to
131 establish the potential fumonisin contamination. All four strains were able to produce fumonisins
132 in peanuts (Table 1). Examples of *A. niger* growth on peanuts are given in Supporting Information
133 1.

134 Table 1: Production of Fumonisin by *Aspergillus niger* in Peanuts after 7-14 Days Growth at 25 °C

Incubation period	IBT 18741 (FB ₂ , mg/kg)		IBT 28753 (FB ₂ , mg/kg)		IBT 29019 (FB ₂ , mg/kg)		IBT 29331 (FB ₂ , mg/kg)	
	#1	#2	#1	#2	#1	#2	#1	#2
7 days	5.8	4.8	3.5	1.6	0.43	0.014	2.5	1.0
14 days	9.6	9.1	2.5	3.9	0.56	3.4	3.9	7.6

135

136 The mean production of fumonisin B₂ varied more than 40 fold from 0.22 mg/kg (IBT 29019) to 9.4
 137 mg/kg (IBT 18741). Selected transitions from the LC-MS/MS analysis of infected and spiked
 138 peanuts are shown in the Supporting Information 2. The measured fumonisin concentrations
 139 (**Table 1**) are similar, compared to those detected in grapes and raisins (0.2-8 mg/kg)^{19,20}.

140

141 **Peanut and peanut butter samples**

142 Since *A. niger* produced fumonisins in high contaminations when growing on the peanuts we
 143 decided to screen peanut retail products. It was further decided to focus on peanut butter and
 144 peanuts purchased directly from local markets, as we assumed the quality would be lower than
 145 from the roasted and salted peanuts sold for snacks.

146 Analysis of 15 peanut samples (from local Thai markets) and 9 peanut butter brands (peanuts
 147 originating from 4 countries) all analyzed in triplicate (as well as a lot of samples during method
 148 development and validation) showed no indications of fumonisins (LOD: 0.004 mg/kg). Even
 149 though *A. niger* is commonly isolated from peanuts⁹⁻¹⁴, and that when it is grown on the peanuts,
 150 fumonisins are produced in high quantities (up to 2000 higher than our detection limit), we
 151 suppose that the combination of: i) extensive damage to the nuts made by *A. niger* (in some cases
 152 they are heavily digested, see supporting information), and: ii) the high attention to reduce
 153 aflatoxin exposure¹⁷ effectively removes *A. niger* infected peanuts and thus fumonisins.

154 Altogether these results indicate that fumonisin exposure in peanuts is not health problem.

155

156 **Supporting Information:**

157 1: Example pictures of *A. niger* growth on peanuts.

158 2: LC-MS/MS chromatograms of spiked and non-spiked peanut samples.

159 Supporting information is available free of charge via the Internet at <http://pubs.acs.org>.

160

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163

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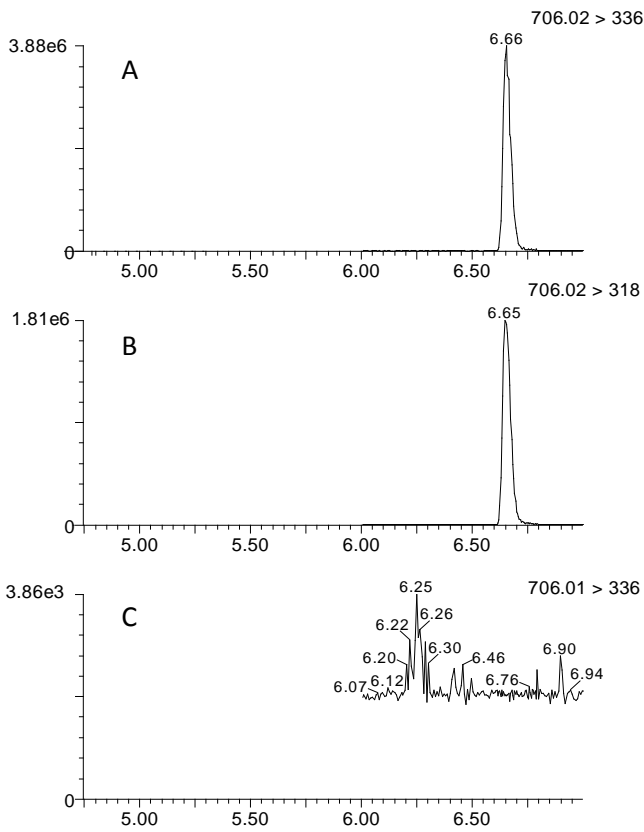
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235
236

Supporting information 1: Examples of *A. niger* growth on peanuts.



Supporting information 2:

MS/MS chromatograms of spiked and blank peanut samples. A: Qualifier of a spiked peanut sample (4 mg/kg). B: Quantifier of a spiked peanut sample (4 mg/kg). C: Qualifier of a peanut sample.



Paper 7

”Single kernel analysis of fumonisins and other fungal metabolites in maize from South African subsistence farmers”

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Single-kernel analysis of fumonisins and other fungal metabolites in maize from South African subsistence farmers

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Fumonisin are important *Fusarium* mycotoxins mainly found in maize and derived products. This study analysed maize from five subsistence farmers in the former Transkei region of South Africa. Farmers had sorted kernels into good and mouldy quality. A total of 400 kernels from 10 batches were analysed; of these 100 were visually characterised as uninfected and 300 as infected. Of the 400 kernels, 15% were contaminated with 1.84–1428 mg kg⁻¹ fumonisins, and 4% ($n = 15$) had a fumonisin content above 100 mg kg⁻¹. None of the visually uninfected maize had detectable amounts of fumonisins. The total fumonisin concentration was 0.28–1.1 mg kg⁻¹ for good-quality batches and 0.03–6.2 mg kg⁻¹ for mouldy-quality batches. The high fumonisin content in the batches was apparently caused by a small number (4%) of highly contaminated kernels, and removal of these reduced the average fumonisin content by 71%. Of the 400 kernels, 80 were screened for 186 microbial metabolites by liquid chromatography-tandem mass spectrometry, detecting 17 other fungal metabolites, including fusaric acid, equisetin, fusaproliferin, beauvericin, cyclosporins, agroclavine, chanoclavine, rugulosin and emodin. Fusaric acid in samples without fumonisins indicated the possibility of using non-toxinogenic *Fusaria* as biocontrol agents to reduce fumonisin exposure, as done for *Aspergillus flavus*. This is the first report of mycotoxin profiling in single naturally infected maize kernels.

Keywords: cereals; fumonisins; LC/MS; mycology

Introduction

Fumonisin are a group of naturally occurring, polyketide-derived mycotoxins produced mainly by *Fusarium verticillioides* and *Fusarium proliferatum* (Rheeder et al. 2002). Recently, fumonisins B₂ and B₄ (FB₂, FB₄) were also detected in *Aspergillus niger* (Frisvad, Smedsgaard et al. 2007; Noonim et al. 2009) and *Tolypocladium* (Mogensen et al. 2011). Fumonisin production has also been detected in a single strain of *Alternaria arborescens* (as *Alternaria alternata* f. sp. *lycopersici*) (Chen et al. 1992) that has since lost this ability (Solfrizzo et al. 2005). Fumonisin are an important health risk because they are potentially carcinogenic and cause toxicoses in humans and animals (Sydenham et al. 1991). The effects of fumonisins include outbreaks of equine leukoencephalomalacia, and porcine pulmonary oedema and hydrothorax (Harrison et al. 1990; Thiel et al. 1991).

Fusarium species occur worldwide in maize, where they infect the cob during flowering. They can produce high amounts of fumonisins in tropical and subtropical

regions (Shephard et al. 1996; Marasas 2001; Reddy et al. 2009) whereas in colder regions the fumonisin contamination is much lower (Logrieco et al. 2002; Miller 2008). In addition to maize-derived products, fumonisins have also been detected in rice (Yazar and Omurtag 2008), black tea leaves (Martins et al. 2001), asparagus (Logrieco et al. 1998) and pine nuts (Marin et al. 2007). Recently *Aspergillus niger* has been shown to be the source of fumonisins in wine and raisins (Knudsen et al. 2010; Logrieco et al. 2010; Mogensen, Larsen et al. 2010). Because of their ubiquitous natural occurrence in maize and maize-based foods (Shephard et al. 1996), fumonisins have attracted considerable global attention from food safety authorities. Consequently, international consultations and risk assessments have resulted in the United States Food and Drug Administration (USFDA) recommending a maximum level in maize for human consumption of 2–4 mg kg⁻¹ total fumonisins depending on the maize product (USFDA 2001), and the European Union setting a regulatory limit of 0.2–2 mg kg⁻¹ (European Union 2007).

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Maize is the staple cereal food grown and it is consumed by rural farming communities of Africa, especially in the Transkei region in South Africa (Shephard et al. 2002), where the effects of fumonisins were first discovered. The Transkei region has one of the highest incidences of oesophageal cancer in the world, which seems to be associated with fumonisin intake (Makaula et al. 1996; Shephard et al. 2002).

Analysis of single kernels has been performed previously with different methodologies, for instance: GC-MS for ergosterol in wheat kernels (Dong et al. 2006), ELISA for deoxynivalenol in wheat kernels (Sinha and Savard 1997), and DESI-MS for fumonisin in the surface of spiked maize kernels (Maragos and Busman 2010).

In this study we examined subsistence-grown maize from the former Transkei region of South Africa sorted into high- and low-quality cobs by the farmers themselves. We surveyed the fumonisin content of single maize kernels to establish the effects of manual visual sorting and determine the levels of fumonisin in the single kernel. Since numerous kernels appearing infected did not contain fumonisins, a further 20% of the analysed kernels were reanalysed for 186 microbial metabolites by a multi-toxin method to detect compounds from non-fumonisin producing *Fusaria* as well as post-harvest contaminants as *Penicillia* and *Aspergilli*.

Materials and methods

Unless otherwise stated, all solvents were high-performance liquid chromatography (HPLC) grade and other chemicals were analytical grade from Sigma-Aldrich (St. Louis, MO, USA). Water was purified on a Milli-Q system (Millipore, Bedford, MA, USA). Certified standard FB₁, FB₂ and FB₃ were from Biopure (Tulln, Austria). Standards of FB₄ and FB₆ were from previous studies (Månsson et al. 2010).

Maize samples

Ten batches of maize (approximately 500 g) were obtained from subsistence farmers in the Centane magisterial district of the former Transkei region, South Africa. Five batches had been classified by the farmers who grew them as high-quality cobs, good maize to be used for human consumption, and five as mouldy quality (mouldy) maize cobs. The mean percentage of infected/damaged kernels of each batch was determined by counting infected kernels in a random selection of 100.

Isolation and identification of mycoflora

Kernels were surface sterilised in 0.4% sodium hypochlorite for 2 min, then washed in sterile water. Six kernels were placed on each of the following media: dichloran 18% glycerol agar (DG18) (Hocking and Pitt 1980), Czapek iprodione dichloran agar (CZID) (Abildgren et al. 1987), and potato-carrot-manganese agar (PCMA) (Sørensen et al. 2009). In addition, six visibly infected or damaged kernels from each batch were placed on 2% water agar, resulting in an overall analysis of 24 kernels per batch and a total of 240 kernels. Plates were incubated for 7 days at 25°C in darkness. *Fusarium*, *Penicillium* and *Aspergillus* species were sub-cultured onto media appropriate for the genus and identified as previously described (Frisvad and Samson 2004; Samson et al. 2006, 2010; Frisvad, Larsen et al. 2007; Wulff et al. 2010), including secondary metabolite profiling and comparison to ex-type cultures (Nielsen and Smedsgaard 2003).

Extraction of kernels

For each maize batch, kernels were divided into groups of undamaged/uninfected and damaged/infected. Based on visual inspection of each batch, ten uninfected and 30 infected kernels were selected for chemical analysis, for a total of 400 kernels.

A single kernel was transferred into a 5-ml cryotube containing ten steel balls (diameter = 3 mm) and 1.5 ml distilled water added. The cryotubes were shaken for 5 min by a Mini Bead-beater (Biospec Products Inc., Bartlesville, OK, USA). Afterwards, 1.5 ml acetonitrile were added and the tubes were shaken on a shaking platform for 30 min. The mixture was centrifuged at 6000 g and 1.5 ml supernatant filtered through a 0.45-µm polytetrafluoroethylene filter and used directly for analysis.

LC-MS/MS fumonisin analysis

Liquid chromatography tandem mass spectrometry (LC-MS/MS) was performed on a Quattro Ultima triple mass spectrometer (Micromass, Manchester, UK) with an electrospray ionisation (ESI) source as previously described (Mogensen, Frisvad et al. 2010). Separations were performed on a 50-mm Gemini C₆-phenyl column (Phenomenex, Torrance, CA, USA), using an acidic linear gradient from 20% acetonitrile to 55% in 6 min at a flow rate of 0.3 ml min⁻¹. Tandem MS was performed in ESI⁺, with the MS in multiple reaction-monitoring (MRM) mode (Mogensen, Larsen et al. 2010; Mogensen, Frisvad et al. 2010). Transitions were: FB₂ and FB₃ *m/z* 706 → 336 and 706 → 512; FB₄ *m/z* 690 → 320 and 690 → 514; and FB₁ and FB₆ *m/z* 722 → 334 and

722 → 528. A few extracts from kernels with the highest infection rate were additionally screened by LC-time-of-flight (TOF)-MS for other fumonisin analogues (Nielsen and Smedsgaard 2003).

Fumonisin extraction was validated by spiking single kernels of undamaged/uninfected maize. The kernels were split with a scalpel and spiked with fumonisin standards to final individual FB₁ and FB₂ concentrations of 6.7, 2.3, 0.75, 0.25 and 0.081 mg kg⁻¹. The kernels were air dried at room temperature for 1 h before extraction. Experiments were performed in triplicate on 3 different days. The apparent recovery was calculated from acetonitrile:water (1:1)-diluted samples with similar concentrations of fumonisins.

LC-MS/MS multi-mycotoxin screening

From the 400 extracts analysed for fumonisins, 80 were selected from the 10 batches to represent a combination of high, low and no amounts of fumonisins. The 80 samples (eight from visually non-infected kernels and 72 from visibly infected or damaged kernels) were analysed for 186 bacterial and fungal metabolites using a QTrap 4000 LC-MS/MS System (Applied Biosystems, Foster City, CA, USA) equipped with a TurboIonSpray ESI source and an 1100 Series HPLC System (Agilent, Waldbronn, Germany), as described by Vishwanath et al. (2009). Chromatographic separation was performed at 25°C on a 15 cm Gemini C₁₈-column. The chromatographic methods and chromatographic and MS parameters for the investigated analytes were as described by Vishwanath et al. (2009). ESI-MS/MS was performed in time-scheduled MRM mode in both positive and negative polarities in two separate chromatographic runs per sample by scanning two fragmentation reactions per analyte. The MRM detection window of each analyte was set to its expected retention time ±30 s in positive mode and ±60 s in negative mode.

Results and discussion

Fungal infection rate of the maize batches

The good-quality batches all had a rate of infected/damaged kernels below 3% ($n = 100$), while the rate in mouldy batches was 6–19%, demonstrating a clear difference in maize quality (Table 1). The average mass of infected and uninfected kernels was 0.29 g and 0.37 g ($n = 14$ each) with a relative standard deviation (RSD) of 14% and 16%, respectively. The visibly infected or damaged kernels plated on water agar showed a 17–83% *Fusarium* infection rate (Table 1), with only fumonisin-producing *Fusaria*. Primarily only *F. subglutinans* and *F. verticillioides* were found, as previously found in South African maize (Rheeder et al.

1992; van der Westhuizen et al. 2011b). The less heat-tolerant *F. graminearum* was found only once among the randomly selected kernels. *Penicillium concavogulosum* was found in four batches, *A. wentii* in three, *Eurotium* sp. in one, *P. aurantiogriseum* in two, *P. crustosum* in two, *P. pittii* in one and *P. brevicompactum* in one.

Validation of fumonisin extraction

The three calibration curves from different days had R^2 -values of FB₁: 0.9932, 0.9986, 0.9974; and FB₂: 0.9974, 0.9978, 0.9984. The RSDs of the slopes were 7.8% (FB₁) and 16.1% (FB₂). The average RSD of the spiked sample ($n = 3$) for FB₁ was 12% (range = 7.5–19%) and for FB₂ 17% (range = 3.4–27%). The limit of quantification (RSD < 20%) was 0.25 mg kg⁻¹ for FB₁ and 0.081 mg kg⁻¹ for FB₂. The limit of detection (LOD) was estimated to an s/n ratio of 5:1, giving an LOD_{FB1} of 0.16 mg kg⁻¹ and LOD_{FB2} of 0.057 mg kg⁻¹. The average apparent recovery for FB₁ was 121% with an average RSD of 25%, and for FB₂ was 115% with an average RSD of 20%.

Fumonisin content in single maize kernels

When single kernels were analysed for fumonisin, all ten batches contained positive kernels. Of the 400 tested kernels, 59 (15%) were positive for fumonisin. Of these, all were positive for FB₁, FB₂, FB₃ and FB₄, except two that did not contain FB₃ and FB₄. LC-TOF-MS screening of a few highly infected kernels also contained fumonisins A₁, A₂, C₁, C₃ and C₄, although in amounts much lower than FBs (data not shown).

None of the maize kernels that were visually characterised as uninfected contained detectable amounts of fumonisins, whereas 3–40% of the visually infected kernels were contaminated with fumonisins (Table 1). The total fumonisin concentration (FB₁, FB₂, FB₃ and FB₄) of single kernels in all the batches was 1.8–1428 mg kg⁻¹. Interestingly, the few low-quality kernels sorted from the good-quality cob batches generally had a higher fumonisin content compared with the poor-quality maize. However, kernels that were visually infected were more frequently found in the poor-quality cob batches.

Using the infection rate, average kernel weight, batch weight and average fumonisin concentration, the total fumonisin content in the batches was estimated. The estimated total fumonisin content of the good-quality batches was 0.28–1.1 mg kg⁻¹ (average = 0.71 mg kg⁻¹), while batches characterised as mouldy contained 0.03–6.2 mg kg⁻¹ fumonisins (average = 2.9 mg kg⁻¹). This showed a marked

Table 1. Infection rate of batches and fumonisins (FB) content in uninfected and infected maize kernels ($n_{\text{total}} = 400$).

Batch	416	438	440	447	448	404	429	431	432	455
Quality ^a Infection rate (%) ^{b,c}	Good 66	Good 66	Good 83	Good 66	Good 100	Mouldy 100	Mouldy 83	Mouldy 66	Mouldy 83	Mouldy 83
Dominating FB-producing fungi (%) ^{b,d}	<i>F. sub</i> (17)	<i>F. vert</i> (17), <i>F. sub</i> (17)	<i>F. vert</i> (17)	<i>F. vert</i> (50)	<i>F. sub</i> (33), <i>F. vert</i> (17)	<i>F. vert</i> (50), <i>F. sub</i> (33)	<i>F. vert</i> (17)	<i>F. vert</i> (17)	<i>F. sub</i> (33)	<i>F. vert</i> (17)
Other fungi ^{b,c,d}	<i>P. conca</i> , <i>A. wentii</i> , <i>P. auran</i>	<i>P. conca</i> , <i>P. auran</i>	<i>P. conca</i>	<i>A. wentii</i>		<i>Eurotium</i> sp.		<i>P. crust</i> , <i>A. wentii</i>	<i>P. pittii</i> , <i>P. brevi</i>	<i>P. conca</i> , <i>P. crust</i>
Visibly infected or damaged kernels ($n = 100$) (%)	2	3	3	3	3	6	14	7	19	12
Uninfected kernels contaminated with FB ($n = 10$) (%)	0	0	0	0	0	0	0	0	0	0
Visually infected kernels contaminated with FB ($n = 30$) (%)	20	27	27	13	10	23	10	40	23	3
FB contents in infected kernels (minimum – maximum, mg kg ⁻¹)	34.4–1428	4.0–90	1.9–715	1.8–432	5.8–929	2.1–257	39–736	3.2–968	4.6–713	8.4
Mean FB content in infected kernels (mg kg ⁻¹) ^e	330	49	114	149	328	54.8	297	166	159	8.4
Median FB in infected kernels (mg kg ⁻¹)	53	42	30	82	50	4.8	117	33	41	8.4
Total FB content (mg kg ⁻¹) ^f	1.1	0.28	0.79	0.52	0.85	0.67	3.7	4.1	6.2	0.03

Notes: Fumonisin content is the total FB₁, FB₂, FB₃ and FB₄ content.

F. vert = *F. verticillioideus*; *F. sub* = *F. subglutinans*; *P. auran* = *P. auranitiogriseum*; *P. brevi* = *P. brevicompactum*; *P. conca* = *P. concavurugulosum*; *P. crust* = *P. crustosum*.

^aAs sorted by the five subsistence farmers.

^bSurface sterilisation in hypochlorite prior to plating for a total of 180 randomly selected kernels (18 per batch with six on DG18, six on CZID and six on PCMA).

^cPlating of 60 visibly infected or damaged kernels on water agar per batch.

^dNot including sterile mycelia and species found only once among the 240 kernels.

^eLC-MS/MS on Ultima QQQ. LOD_{FB1} of 0.16 mg kg⁻¹ and LOD_{FB2} of 0.057 mg kg⁻¹.

^fTotal FB content was calculated from the infection rate, mean kernel weight, batch weight and mean fumonisin concentration.

difference in the fumonisin content of the high- and low-quality cob batches.

Of the ten batches, two exceeded the maximum level of 4 mg kg^{-1} recommended by the USFDA in maize for human consumption (USFDA 2001), and four exceeded the European Union regulatory limit of 1 mg kg^{-1} (European Union 2007). The values for fumonisin were similar to previous findings in South African subsistence maize (van der Westhuizen et al. 2008, 2010a). In a previous study van der Westhuizen et al. (2008) found levels of fumonisins in the mouldy maize (as classified on cobs by the farmers) that were higher (mean levels in different areas over two seasons $4.85\text{--}12.9\text{ mg kg}^{-1}$) than those found in this study.

Multi-mycotoxin screening of selected kernels

In the previous analysis, the kernels were only tested for fumonisin, however since only 15% of the kernels were contaminated with fumonisin, we performed a multi-mycotoxin screening for identification of other toxins from non-fumonisin producing fungi. As well as to investigate the presence of compounds produced by post-harvest fungi, e.g. *P. concavorugulosum* and *A. wentii*.

Eighty kernels were selected and analysed by the multi-detection method for 186 bacterial and fungal metabolites, as described by Vishwanath et al. (2009). The results are shown in Table 2.

Fumonisin was detected in 50 of the 80 samples (Table 2), and only in four of the samples were no microbial metabolites detected. Other frequently detected metabolites were fusaric acid ($n=53$), emodin ($n=27$), chanoclavine ($n=25$), equisetin ($n=14$) and fusaproliferin ($n=12$). Metabolites found in a low number of samples were: agroclavine ($n=3$), altertoxin-I ($n=3$), beauvericin ($n=7$), moniliformin ($n=4$), nivalenol ($n=4$), rugulosin ($n=4$), and tentoxin ($n=3$). Other metabolites found in single cases were: alternariol, calphostin C, and cyclosporins A, C and H. To our knowledge, this is the first report of the fungal metabolites agroclavine, calphostin C, chanoclavine, cyclosporins, equisetin, rugulosin and tentoxin, as well as the bacterial metabolites nonactin, monactin and valinomycin in maize.

Table 3 shows a summary of Table 2 with the suspected producers of the metabolites as supported by mycological examination as well as if the compounds are produced pre- or post-harvest. Of the detected compounds and mycotoxins, many are presumably produced pre-harvest based on the natural field growth patterns of *Alternaria* and *Fusarium*. However, a lower number of toxins, generally considered to be produced post-harvest, were also detected, mostly those originating from *Penicillium*, *Aspergillus* and *Eurotium*. The finding of pre-harvest mycotoxins occurring from

several *Fusaria* in the same maize kernels points toward the co-existence of the producers rather than species succession which could explain the remaining compounds originating from several species and genera.

Beauvericin, fusaproliferin and moniliformin (Table 1) are metabolites of *F. subglutinans*. Kernels with fusaproliferin and/or moniliformin often showed lower amounts of fumonisins, which could indicate the presence of *F. subglutinans*, which is known to be a poor fumonisin producer (Sala et al. 1994). Furthermore, samples containing fumonisin and fusaric acid had an average fusaric acid content of 37 mg kg^{-1} , while visually infected samples without fumonisin had a significantly higher average content of 112 mg kg^{-1} . Fusaric acid is a pre-harvest compound, a plant growth inhibitor and is suggested to be antifungal and antineoplastic (Bacon et al. 1996; Diniz and Oliveira 2009). Its presence in samples without fumonisins strongly indicates that non-toxinogenic *Fusaria* were present in the maize, and supports the possibility of using these fungi as biocontrol agents against fumonisin-producing species, as is the case for *A. flavus* (Antilla and Cotty 2002; Cleveland et al. 2003; Abbas et al. 2006).

Equisetin is known to be produced by *F. equiseti*, which also produces nivalenol; however, this species was not detected. Nivalenol could also originate from *F. graminearum*, but this species was found only on a single kernel. However, the mycological screening was conducted on a limited number of kernels, so detection and isolation of a few *F. equiseti* or *F. graminearum* in samples dominated by *F. verticillioides* and *F. subglutinans* would have been difficult. In addition, different kernels were used for the mycological examination and for chemical analysis.

The cyclosporins could have originated from either *F. solani* or *Tolypocladium niveum*, both previously described as producers of cyclosporins (Dreyfuss et al. 1976; Isaac et al. 1990; Proksa et al. 1991; Sugiura et al. 1999).

Several *Alternaria* species can produce alternariol, tentoxin and altertoxin I, and the finding of these compounds was expected since *Alternaria* spp. grow superficially on many grain types (Andersen et al. 2002). Agroclavine, chanoclavine and rugulosin indicate that *P. concavorugulosum* (a known producer of these compounds; Frisvad 1989) was present in batches 429, 432, 438 and 440, which was validated by the mycological findings of this species these batches. Emodin is a common metabolite of many *Eurotium* species and *A. wentii* (Samson et al. 2010), which are common superficial contaminants on dried grain and were found on some kernels from two good and two mouldy batches. The presence of emodin and chanoclavine in the visually uninfected kernels was not surprising, as growth is superficial and thus easily

Table 2. Fungal and bacterial metabolites in single maize kernels by multi-detection MS/MS analysis.

Batch	Maize quality ^a	Sample number	FB ₁ (mg kg ⁻¹)	FB ₂ (mg kg ⁻¹)	FB ₃ (mg kg ⁻¹)	FB ₄ (mg kg ⁻¹)	Fusaric acid (mg kg ⁻¹)	Equisetin (µg kg ⁻¹)	Chanoclavine (µg kg ⁻¹)	Emodin (µg kg ⁻¹)	Other compounds
416	Good	2486	3.03	0.53	0.38	0.17	1.9	—	—	—	
		2489	—	—	—	—	—	—	4.6	—	
		2491	94	39	8.5	8.8	8.1	68	—	—	
		2502	1910	866	167	127	7.3	55	—	—	Beauvericin, nivalenol
		2506	747	246	71	55	17	35	0.78	—	Beauvericin, nivalenol
438	Good	2527	9.3	2.9	1.1	0.57	17	—	0.53	—	
		2529	46	43	1.3	4.4	16	—	83	6.1	Agroclavine, rugulosin
		2531	21	8	2.9	2	12	—	—	9.2	Fusaproliferin, alternariol
		2533	—	0.14	—	—	46	—	—	18	Fusaproliferin
		2535	—	—	—	—	212	14	15	—	Fusaproliferin, monactin, nonactin, valinomycin
		2543	0.07	—	—	—	125	—	—	3.7	Altartoxin-I
		2545	—	—	—	—	—	7756	—	—	Fusaproliferin, beauvericin, nivalenol, tentoxin
440	Good	2546	149	26	6.8	4	26	—	0.52	—	
		2550	48	16	7.8	4.4	24	—	—	—	
		2558	—	—	—	—	0.2	—	—	7.3	
		2566	—	—	—	—	—	—	—	—	
		2567	0.18	0.062	—	—	—	—	1.3	—	
		2582	794	221	43	31	31	218	5.6	6.8	Rugulosin
		2584	4.7	1.2	0.61	0.34	63	—	6	—	
		2586	—	—	—	—	5.1	—	0.82	—	
		2591	37	11	4.2	2.9	12	—	—	35	
		2592	34	13	2.9	2.2	2.8	—	—	—	
		2603	—	0.033	—	—	—	—	—	—	
		2619	2.6	0.76	0.59	0.43	24	—	—	—	
447	Good	2622	—	—	—	—	—	—	—	—	
		2626	—	—	—	—	—	—	—	—	
		2629	730	203	47	26	1.7	—	—	46	Beauvericin
		2632	—	—	—	—	1.2	—	—	9.7	
		2633	—	—	—	—	—	—	—	—	
		2644	—	—	—	—	—	—	—	12	Beauvericin, tentoxin, altartoxin-I
		2647	5.6	0.26	0.6	—	4.7	1691	—	—	Moniliformin
		2649	1137	481	84	74	38	—	1	—	
404	Mouldy	2652	0.58	0.08	—	—	—	—	5.9	3.9	Fusaproliferin, moniliformin
		2659	—	—	—	—	165	—	—	—	Fusaproliferin
		2665	—	—	—	—	73	12	—	—	
		2681	—	—	—	—	—	1939	1.1	10	Cyclosporins A, C, H
		2687	206	72	13	11	25	—	3.8	—	
		2696	3	0.74	0.28	—	8.1	—	—	—	
		2699	1.8	0.57	0.2	0.0058	—	—	—	—	Fusaproliferin
		2704	—	—	—	—	40	—	—	—	

(continued)

Table 2. Continued.

Batch	Maize quality ^a	Sample number	FB ₁ (mg kg ⁻¹)	FB ₂ (mg kg ⁻¹)	FB ₃ (mg kg ⁻¹)	FB ₄ (mg kg ⁻¹)	Fusaric acid (mg kg ⁻¹)	Equisetin (µg kg ⁻¹)	Chanoclavine (µg kg ⁻¹)	Emodin (µg kg ⁻¹)	Other compounds
429	Mouldy	2706	—	0.038	—	—	—	—	—	—	
		2708	78	32	12	6.4	2.4	—	—	—	
		2720	—	—	—	—	—	—	0.63	14	
		2727	—	—	—	—	1.0	—	—	36	
		2728	—	—	—	—	—	—	2.1	—	
		2730	886	385	20	20	4.4	—	—	—	
		2738	158	27	7.1	3.4	26	—	—	17	Rugulosin
		2741	—	—	—	—	—	—	—	16	
		2745	—	—	—	—	21	—	—	—	
		2758	—	—	—	—	—	33	126	4.4	Agroclavine
431	Mouldy	2766	1277	361	47	21	0.2	—	1.1	—	
		2767	2.4	0.64	0.25	0.0076	16	—	—	—	
		2768	8.5	1.9	0.61	0.39	0.4	—	0.59	—	
		2776	—	—	—	—	—	—	2.5	6.2	Fusaproliferin
		2781	—	—	—	—	23	24	1.5	—	
		2782	265	39	31	14	4.0	—	13	3.3	Fusaproliferin
		2784	932	254	47	34	55	—	—	—	Beauvericin, moniliformin
		2792	51	14	6.7	3.9	8.4	—	0.04	—	
		2798	—	—	—	—	—	—	—	—	Nivalenol
		2807	0.43	0.13	—	—	0.8	—	106	75	Fusaproliferin, agroclavine, altertotoxin-I, rugulosin
432	Mouldy	2810	—	—	—	—	251	—	—	5.2	
		2815	—	0.041	—	—	—	—	—	—	
		2817	—	—	—	—	—	101	—	—	Tentoxin
		2820	30	22	4.6	9	23	—	—	—	
		2821	15	9	1.5	2.1	6.6	—	—	9.5	
		2822	—	—	—	—	—	—	—	14	Moniliformin
		2825	—	—	—	—	—	—	—	—	Calphostin C
		2826	739	306	83	92	13	24	—	—	Beauvericin
		2828	226	90	23	25	22	—	—	—	
		2829	3.5	2.1	0.31	0.32	17	—	—	—	
455	Mouldy	2833	4	2.9	0.61	1.3	2.0	—	—	—	
		2839	—	—	—	—	—	—	—	3.5	
		2847	0.47	0.35	0.14	0.18	0.6	—	—	8.9	
		2855	—	—	—	—	—	—	0.46	—	
		2856	1.1	0.16	—	—	49	—	—	—	
		2858	0.1	0.056	—	—	—	—	—	—	
		2860	5.9	1.2	0.74	0.35	29	—	—	—	
		2865	—	0.031	—	—	27	—	—	169	
		2868	—	0.036	—	—	—	709	—	58	Fusaproliferin
		2872	—	0.041	—	—	189	—	—	—	

Notes: '—' indicates that the levels were below the level of detection.
^aAs sorted by the five subsistence farmers.

Table 3. Distribution and likely producer of metabolites found in visually infected and uninfected kernels.

Mycotoxin	Percentage in uninfected kernels (n = 8)	Percentage in infected kernels (n = 72)	Likely producer	Supported by microbiology	Produced pre- or post-harvest
Alternariol	0	1	<i>Alternaria</i>	No	Pre ^a
Altartoxin-I	13	3	<i>Alternaria</i>	No	Pre ^a
Beauvericin	13	8	<i>F. subglutinans</i>	Yes	Pre
Calphostin C	0	1	<i>Cladosporium</i>	No	Pre ^a
Cyclosporin A, C and H	13	0	<i>F. solani</i> or <i>Tolypocladium</i>	No	Pre
Equisetin	25	17	<i>F. equiseti</i>	No	Pre
Fumonisin B ₁	0	60	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fumonisin B ₂	13	65	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fumonisin B ₃	0	51	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fumonisin B ₄	0	49	<i>F. verticillioides</i> , <i>F. subglutinans</i>	Yes	Pre
Fusaproliferin	0	17	<i>F. subglutinans</i>	Yes	Pre
Fusaric acid	13	64	<i>F. verticillioides</i> , <i>F. subglutinans</i> and other related	Yes	Pre
Moniliformin	0	6	<i>F. subglutinans</i>	Yes	Pre
Nivalenol	13	4	<i>F. equiseti</i> , <i>F. graminearum</i>	(Yes) ^b	Pre
Tentoxin	13	3	<i>Alternaria</i>	No	Pre ^a
Agroclavin	13	3	<i>P. concavurugulosum</i>	Yes	Post
Chanoclavine	38	31	<i>P. concavurugulosum</i>	Yes	Post
Emodin	75	29	<i>Eurotium</i> , <i>A. wentii</i>	Yes	Post
Rugulosin	0	6	<i>P. concavurugulosum</i>	Yes	Post

Notes: ^aAs superficial growth usually close to harvest.

^b*F. graminearum* was only found on a single kernel.

abraded off the kernels so they appear uninfected. The bacterial metabolites monactin, nonactin and valinomycin were found only once, and in the same sample.

Even though aflatoxins, like fumonisins, are considered important mycotoxins in African maize, no aflatoxins were found in any of the analysed samples (Shephard 2008). These findings corresponded to the fungal isolation results, which found neither *A. flavus* nor *A. parasiticus* from any of the 240 kernels surveyed and also to a previous study conducted in the former Transkei region which also did not find any aflatoxin or *A. flavus* (van der Westhuizen et al. 2011a).

This is the first study to examine the mycotoxin profile of single maize kernels. The maize mycotoxin profiles were diverse, and the detection of metabolites from several fungi on single kernels supports the fact that several fungal species can co-occur on a single kernel. The results showed a low percentage of maize kernels containing fumonisins (15%), and an even lower percentage (<4%) with a high concentration of fumonisins (>100 mg kg⁻¹). The results also confirm that *Fusarium* mycotoxins are non-homogeneously distributed in maize (van der Westhuizen et al. 2011a). Although fumonisins appeared to be the cause of the most serious mycotoxin problem since they were the most widespread of the analysed mycotoxins, the fumonisin contamination in the maize samples was mainly caused by a low percentage of

highly infected kernels. This supports the approach if a more thorough sorting of the subsistence grown maize is performed to decrease the fumonisin concentration. A theoretical calculation of the effect of removing the highly infected kernels (4%) showed that the average fumonisin concentration decreased by 71% (data not shown) after a simple sorting. This is a substantial effect gained using a very simple method. The strategy of sorting out visibly infected kernels has recently been successfully applied in an intervention study in the same rural Transkei area (van der Westhuizen et al. 2011b). A group of farmers were trained to remove infected kernels from their good-quality maize and achieved a mean fumonisin reduction of 84% by removing a mean of 3.9% by weight, which is very close to our estimated reduction (van der Westhuizen et al. 2010b, 2011a, b).

A valuable addition to these results could be to investigate the mycotoxin chemistry in single kernels, while simultaneously conducting an investigation of the mycology of the same kernel, for a better understanding of how species co-exist and how they are spread across the cob. This could also determine whether several fungal species infect the same kernel or if kernels are infected with only one species. Furthermore, data on single-kernel mycotoxin content may help in designing sampling plans for detecting mycotoxins in large lots. For example, Whitaker et al. (1972) suggested using a negative binominal

distribution and Vargas et al. (2005) showed that approximately six coffee beans per 10,000 beans are contaminated if a lot contains $5 \mu\text{g kg}^{-1}$ of ochratoxin A. The results of our single-kernel mycotoxin analysis give an in-depth view of the extent of mycotoxin-producing fungi in South African Transkei subsistence maize, particularly fumonisins. These findings are valuable for designing sampling plans for large maize lots.

The results support the visual inspection method currently used by the maize-producing farmers and support additional sorting for even more effective removal of mycotoxin-contaminated kernels. Additional mycotoxin reduction might be achieved through biocontrol measures using a non-fumonisin-producing *Fusarium* strain. Finally, the single-kernel approach is promising for the detection of new fungal compounds in maize, as the concentrations of fungal compounds are much higher in the few infected kernels.

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Paper 8

“Production of Fumonisins B₂ and B₄ in *Tolypocladium* species”

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Production of fumonisins B₂ and B₄ in *Tolypocladium* species

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Abstract *Tolypocladium inflatum* is known primarily for its production of the cyclosporines that are used as an immunosuppressive drug. However, we report here the production of the carcinogenic fumonisins B₂ and B₄ by this biotechnologically relevant fungal genus. These mycotoxins were detected in 11 strains tested from three species: *Tolypocladium inflatum*, *T. cylindrosporum*, and *T. geodes*. Production of fumonisins by *Fusarium* spp. and *Aspergillus niger* is highly medium- and temperature-dependent, so the effect of these parameters on fumonisin production by three *T. inflatum* strains was studied. Maximum production was achieved on media with high sugar content incubated at 25–30°C. Since these results demonstrate that fumonisin production could be widespread

within the genus *Tolypocladium*, the potential contamination of commercial cyclosporine preparations with fumonisins needs to be investigated.

Keywords Fumonisin · Cyclosporin · Mycotoxin · *Tolypocladium inflatum* · *Elaphocordyceps subsessilis*

Introduction

Species in the fungal genus *Tolypocladium* are found worldwide as soil-borne insect pathogens and saprotrophs [6, 35]. *Tolypocladium inflatum* W. Gams (syn. *T. niveum* [Rostr.] Bissett) is the best-known species, because it produces the cyclosporines that are used worldwide as an immunosuppressive drug in organ transplant recipients. In addition, the species produces antifungal and insecticidal efrapeptins [4, 16, 18]. *T. inflatum*, an anamorph of *Elaphocordyceps subsessilis* [Petch] G. H. Sung, J. M. Sung and Spatafora, is currently classified within *Ascomycota*, *Sordariomycetes*, *Hypocreales*, *Ophiocordycipitaceae* [41].

Fumonisin (Fig. 1) are regulated mycotoxins that cause human and animal toxicoses when consumed via contaminated maize-based food and feeds [46]. Fumonisin B₁ (FB₁) has furthermore been implicated as a risk factor for neural tube defects in embryos [21, 45]. Fumonisin were first discovered in *Fusarium verticillioides* (Sacc.) Nirenberg [5] and are a significant problem in maize-based products [14, 36, 39, 42]. Fumonisin are also occasionally found in rice [31], black tea leaves [23], asparagus [19], pine nuts [22], and wine [25]. The US Food and Drug Administration recommends that maize should not be used for human consumption at levels above 2–4 ppm total fumonisin [43] while the European Union (EU) has a regulatory limit of 0.2–2 ppm in maize products [10]. *Fusarium*

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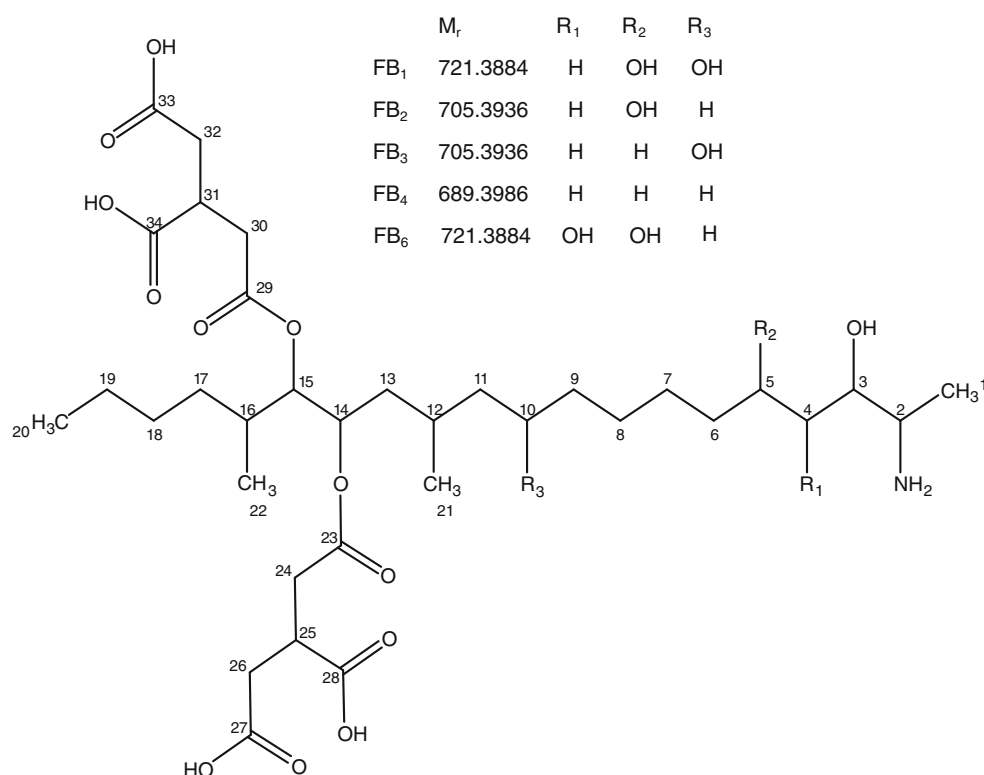
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Fig. 1 Structure of fumonisins B₁, B₂, B₃, B₄ and B₆



is mainly the genus associated with fumonisin production, but a homolog to the *F. verticillioides* fumonisin gene cluster has been found in *Aspergillus niger* Tiegh. [1, 13, 20, 30, 32] and in a single strain of *Alternaria arborescens* E.G. Simmons (referred to as *Alternaria alternata* (Fr.) Kiessl. f. sp. *lycopersici* Grogan, Kimble & Misaghi in [47]). However, of the two only *A. niger* is able to produce fumonisins [13, 20, 30, 37].

During a routine screening of cyclosporine-producing fungi with the multi-detection method described by Vishwanath et al. [44], we were surprised to discover that a *Tolypocladium* strain was able to produce fumonisins. Since this has a potentially important impact on drug safety and genetic analysis, this study was initiated to survey the distribution of fumonisin production in *Tolypocladium* and to examine the influence of media and temperature on fumonisin production by species of this genus.

Materials and methods

Unless otherwise stated all solvents were HPLC grade, chemicals were analytical grade, and water was purified on a Milli-Q system (Millipore, Bedford, MA). Certified reference standards of fumonisin (B₁, B₂, and B₃) were from Romer Labs (Tulln, Austria), and FB₄ and FB₆ were purified from *A. niger* and validated by nuclear magnetic

resonance (NMR) spectroscopy at the Technical University of Denmark [20].

Fungal strains, media, and incubation

Eleven *Tolypocladium* strains and one *Aspergillus niger* strain (IBT 28144 = CBS 101705, NRRL 567, ITEM 7097) were used in this study (see Table 1). To analyze the effect of media on the production of fumonisins ten media were used: Czapek yeast autolysate agar (CYA) [12], Czapek yeast autolysate agar + 5% NaCl (CYAS) [12], dichloran Rose bengal yeast extract sucrose agar (DRYES) [11], potato carrot agar (PCA) [37], malt extract agar (MEA according to Blakeslee) [34], oatmeal agar (OAT) [12], potato dextrose agar (PDA, Difco), V8-juice agar with antibiotics (V8) [37], yeast extract sucrose agar (YES) [12], and dichloran 18% glycerol agar (DG18) [15]. The composition of the media is given in Supplementary Table 1. Media were prepared in 9-cm Petri dishes, each with 17 mL medium, and strains were inoculated as three-point inoculations. Petri dishes were incubated in micro-perforated plastic bags at 25°C in the dark. All samples were analyzed in triplicate on three individual plates. Colony diameter measurements are given as averages of 9 colonies on 3 plates. To analyze the effect of temperature on the production of fumonisins in *T. inflatum*, three strains (IBT 41581, IBT 41582 and IBT 41583) were inoculated

Table 1 Production of fumonisins B₂ (FB₂) and B₄ (FB₄) by *Tolypocladium cylindrosporum*, *T. geodes*, and *T. inflatum*

Species	Strain number	FB ₂	FB ₄
<i>T. cylindrosporum</i>	CCF 1450	+	+
	CCF 2531	+	+
	CCF 3237	+	+
<i>T. geodes</i>	CCF 2548	+	–
	CCF 2579	+	+
	CCF 3299	+	–
<i>T. inflatum</i>	DSM 915	+	+
	DSM 63544	+	+
	IBT 41581	+	+
	IBT 41582	+	+
	IBT 41583	+	+

CCF Culture Collection of Fungi (Prague, Czech Republic); DSM Deutsche Sammlung von Mikroorganismen und Zellkulturen (Braunschweig, Germany); IBT Culture collection at Center for Microbial Biotechnology, DTU Systems Biology (Kgs. Lyngby, Denmark)

on YES and incubated at 15, 20, 25, 30, or 37°C in the dark for 7 days.

Metabolite extraction

Fumonisin were extracted by using the method previously described by Frisvad et al. [13]. Six plugs (6 mm diameter) were cut across the colony diameter through the center. If colony diameters were smaller than 20 mm, plugs were collected from two or three colonies from the plate. The six plugs were transferred to a 2-mL vial, 800 µL of methanol/water (3:1 v/v) was added, and vials were placed in an ultrasonication bath for 1 h. All extracts were filtered through a PTFE 0.45-µm syringe filter (National Scientific, Rockwood, Tennessee) into a new vial and used directly for analysis. Samples were made in triplicate from individual plates. This method was validated as in our previous reports with similar recovery of 75–85% [26, 28].

LC–MS/MS analysis of fumonisins

Fumonisin were analyzed by using liquid chromatography tandem mass spectrometry (LC–MS/MS) as previously described [24] with minor changes. The LC–MS/MS analyses were performed on a Quattro Ultima triple mass spectrometer (Micromass, Manchester, UK) with electrospray ionization (ESI) source, and separations performed on a 50 × 2 mm i.d., 3-µm Gemini C₆-phenyl column (Phenomenex, Torrance, California). A linear gradient was performed from 20% acetonitrile in water with 20 mM formic acid to 55% acetonitrile for 6 min at 0.3 mL/min, increasing to 100% acetonitrile in 30 s at 0.5 mL/min; this

was maintained for 3.5 min before returning to the starting conditions in 6 min. Tandem MS was performed in ESI⁺ with MS operated in multiple reaction monitoring (MRM) mode. Transitions were as previously described [24, 25]; fumonisins B₂ and B₃ quantifier m/z 706→336, qualifier m/z 706→512 a; fumonisin B₄ quantifier m/z 690→320, qualifier m/z 690→514 a; fumonisins B₁ and B₆ quantifier m/z 722→334, qualifier m/z 722→528 a. Reference standards of FB₁–FB₄ and FB₆ were co-analyzed with each analysis sequence. Strains of *T. cylindrosporum* and *T. geodes* (CCF and DSM) were analyzed according to Vishwanath et al. [44] on a similar instrument.

LC–TOF–MS screening

Selected extracts were additionally tested for other fumonisins, efrapeptins, and cyclosporines by liquid chromatography time-of-flight mass spectrometry (LC–TOF–MS) as described in Nielsen et al. [28]. This was performed on an LC system coupled to an orthogonal TOF mass spectrometer (Micromass LCT, Manchester, UK) equipped with an electrospray source [28]. The column was a 50 × 2 mm i.d., 3 µm Luna C₁₈ (II) column (Phenomenex, Torrance, California) and a linear gradient was used with all solutions containing 20 mM formic acid, from 15% acetonitrile in water to 100% acetonitrile in 20 min at 300 µL/min; this was maintained for 3.5 min before returning to the starting conditions in 6 min. The MS operated in ESI⁺ at a scan range of m/z 100–2,000. Peaks not matching the compounds in an internal database with approximately 850 reference standards [29] were matched against the 35,500 structures in Antibase 2009 (John Wiley and Sons, Inc, Hoboken, New Jersey). Extracted ion chromatograms (±0.02 amu) of the [M + H]⁺ ions of the fumonisin series A, C, or P were constructed to search specifically for these compounds.

Results and discussion

A screening for fumonisins found that all eleven *Tolypocladium* strains used in this study produced fumonisins B₂ (FB₂) and nine (82%) also produced fumonisins B₄ (FB₄); however, no FB₁, FB₃ or FB₆ was detected in any of the strains (representative chromatograms in Fig. 2). All *T. cylindrosporum* and *T. inflatum* strains and one *T. geodes* strain (CCF 2579) produced both FB₂ and FB₄, while the other two *T. geodes* strains produced only FB₂ (Table 1).

Three strains of *T. inflatum* (IBT 41581, IBT 41582, IBT 41583) were selected for quantitative analysis of fumonisin production on a variety of different media and at different conditions. We tested 10 different media, and found that on each medium at least one *T. inflatum* strain produced FB₂

Fig. 2 LC–MS/MS analyses of fumonisins B₂ and B₄ in *Aspergillus niger* IBT 28144 and *Tolypocladium inflatum* IBT 41581 extracts from cultures incubated at 25°C for 7 days on YES. **a** Qualifier of a diluted certified FB₂ standard. **b** Quantifier of a diluted certified FB₂ standard. **c** Qualifier of FB₂ in a *T. inflatum* extract. **d** Quantifier of FB₂ in a *T. inflatum* extract. **e** Qualifier of FB₄ in an *A. niger* extract. **f** Quantifier of FB₄ in an *A. niger* extract. **g** Qualifier of FB₄ in a *T. inflatum* extract. **h** Quantifier of FB₄ in a *T. inflatum* extract

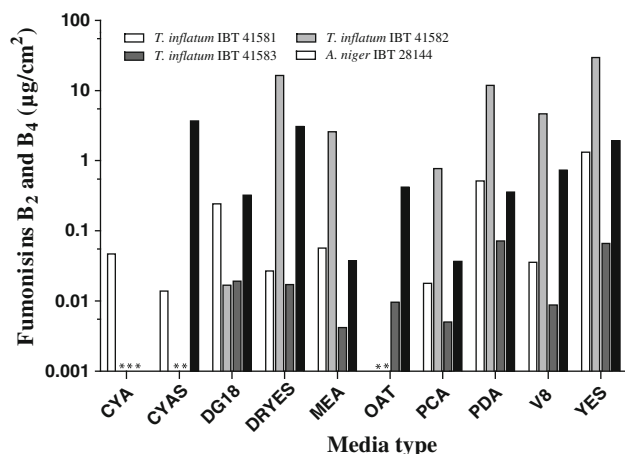
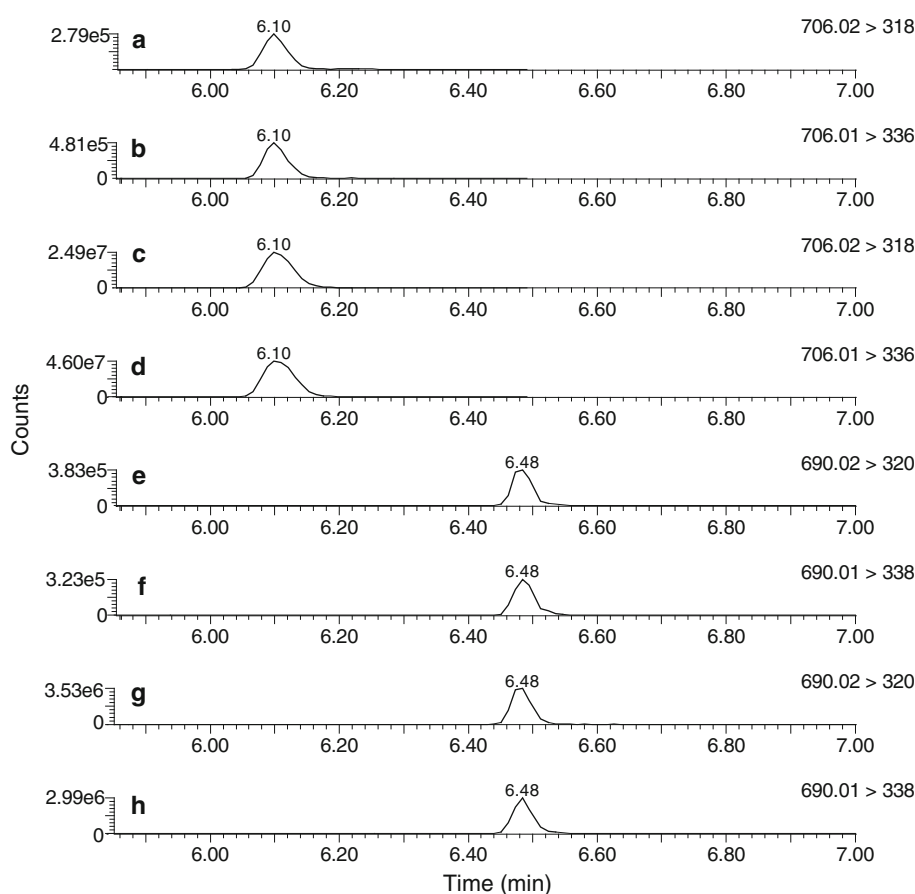


Fig. 3 Production of fumonisins B₂ and B₄ by three *Tolypocladium inflatum* strains (IBT 41581, IBT 41582, IBT 41583) and one *Aspergillus niger* (IBT 28144) on different media at 25°C. *Not detected

and FB₄ (Fig. 3). The maximal production of total fumonisins for the three strains was 1.3 µg/cm² for IBT 41581, 30 µg/cm² for IBT 41582 and 0.072 µg/cm² for IBT 41583. These levels were comparable to both *Fusarium* spp. and *A. niger* in a similar study under similar

conditions, which found fumonisin production of 0.006–22 µg/cm² for *Fusarium* spp. and 0.27–21 µg/cm² for *A. niger* [26]. Relative levels of FB₄ to FB₂ were in the range 0–95%, with most being in the 10–30% range. The media yielding the highest amounts of fumonisins were DRYES, PDA, and YES, while lower amounts were detected in cultures grown on MEA, OAT, CYA, CYAS, PCA, V8, and DG18. *Aspergillus niger* (IBT 28144) produced high amounts of fumonisins on media with high amounts of sugar and salt, e.g., CYAS, DRYES, and YES, with detectable production of fumonisins on the other seven media as shown in Fig. 3. This is in agreement with the results of Frisvad et al. [13]. Fumonisin production by the *T. inflatum* strains was similar to that by both *Fusarium* and *A. niger*, with PDA and YES also supporting substantial fumonisin production [13, 26]. The media used for cyclosporine production often consists of 2–8% carbon source (e.g., sorbose, glucose, or similar) and 1–6% nitrogen source (e.g., peptone, malt extract, or casein acid hydrolysate) with added trace metals [2, 3, 17]. The media most comparable to these in our study are CYA, CYAS, and MEA. On CYA and CYAS, only one strain produced fumonisin, whereas all three strains produced fumonisins on MEA. This suggests that the media used for cyclosporine production might support the simultaneous

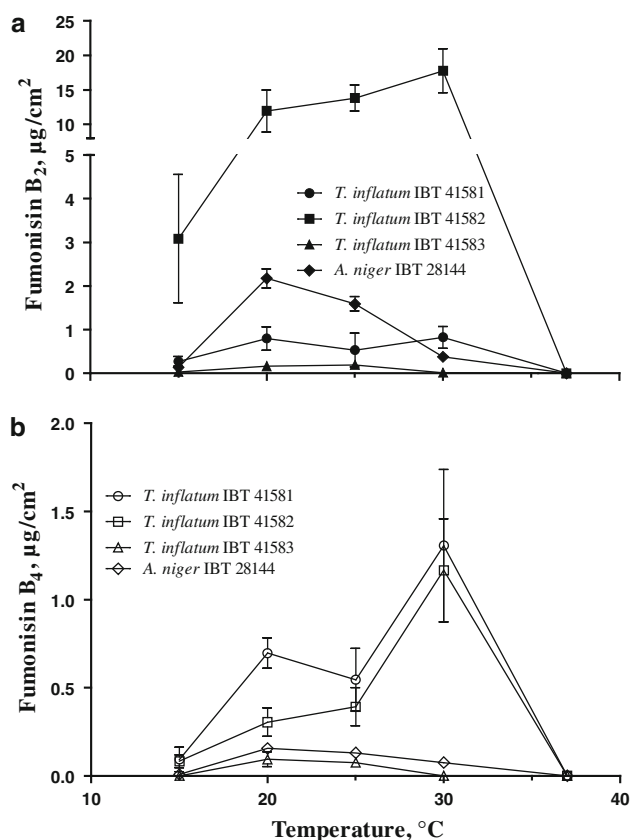


Fig. 4 Production of fumonisins B₂ (a) and B₄ (b) by three strains of *Tolypocladium inflatum* (IBT 41581, IBT 41582 and IBT 41583) and *Aspergillus niger* IBT 28144 incubated for 7 days at 15, 20, 25, 30, and 37 $^{\circ}\text{C}$ on YES. The standard deviation of triplicates is shown by the error bars. Fumonisin concentration was 0 $\mu\text{g}/\text{cm}^2$ at 37 $^{\circ}\text{C}$ because no growth occurred at that temperature

production of fumonisins, although the use of liquid medium compared to solid has to be accounted for.

The three *T. inflatum* strains produced fumonisins in a temperature range of 15–30 $^{\circ}\text{C}$ (Fig. 4). The effect of temperature on fumonisin production was strain-dependent, with two strains giving maximal production at 30 $^{\circ}\text{C}$ and one strain (IBT 41583) showing minimal production at 30 $^{\circ}\text{C}$, and maximum production at 25 $^{\circ}\text{C}$. This is similar to previous observations for *A. niger*, which showed peak fumonisin production at 25–30 $^{\circ}\text{C}$ [26]. Except for one set of experiments in which IBT 41582 produced more FB₄ than FB₂ at 30 $^{\circ}\text{C}$, FB₂ was consistently produced at higher levels than FB₄ (data not shown). All three strains of *T. inflatum* were able to grow at temperatures from 15 to 30 $^{\circ}\text{C}$, with maximal colony diameters after 7 days of incubation at 20 or 25 $^{\circ}\text{C}$ (Fig. 5). No growth was observed at 37 $^{\circ}\text{C}$.

In the three selected *T. inflatum* strains, no traces of fumonisin series A, C, or P were detected by LC–TOF–MS. However, FB₂ and FB₄ were detected in all samples of all

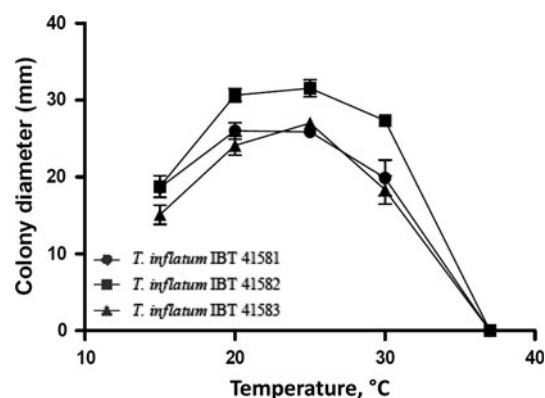


Fig. 5 Colony diameters of three strains of *Tolypocladium inflatum* (IBT 41581, IBT 41582, IBT 41583) after incubation for 7 days at 15, 20, 25, 30, and 37 $^{\circ}\text{C}$ on YES. The standard deviation of triplicates is shown by the error bars

cultures of the three *T. inflatum* strains in repeated experiments, and by the two different techniques of LC–MS/MS and LC–TOF–MS. Fumonisins were detected from cultures grown on different media (PDA, DRYES, and YES), and in separate cultures derived from the same original strain. In addition to FB₂ and FB₄, the three *T. inflatum* strains produced cyclosporines and efraeptins with the fumonisins (data not shown).

This is the first report describing the production of FB₂ and FB₄ in the genus *Tolypocladium*. These findings are relevant because *T. inflatum* is used in the pharmaceutical industry as a producer of cyclosporin A. We recommend that the production strains of cyclosporines are analyzed for production of fumonisins under industrial production conditions to determine if these mycotoxins end up in the final product. Furthermore *Tolypocladium* spp. have been suggested as a possible fungal biological control agent, because of the entomopathogenicity of its production of efraeptins [40]. With the discovery of fumonisin production by *Tolypocladium* spp., however, this raises concerns that these toxic metabolites may enter the environment and the food chain and pose a risk to humans and animals.

To date more than 20 species within the three genera *Fusarium*, *Aspergillus*, and *Tolypocladium* have been shown to produce fumonisins [13, 33]. *Alternaria arborescens* (as *Alternaria alternata* f. sp. *lycopersici*) was reported to produce fumonisins, but the results were later questioned and found to be due to only the chemically similar AAL toxin and not fumonisin [8, 9, 38]. So far, *Fusarium* spp. produce the greatest diversity of fumonisins, namely FB₁–FB₅ and the A, C, and P series [5, 7, 27], whereas *A. niger* produces only FB₂, FB₄, and FB₆ [13, 20, 30]. The discovery reported here shows that *Tolypocladium* spp. have a fumonisin profile similar to *A. niger*, but quite different from *Fusarium* spp. Because of these similar profiles, *Tolypocladium* spp. and *A. niger* could have a

higher degree of similarity in their fumonisin gene cluster compared to *Fusarium* spp., but this needs to be established by genome sequencing of the strains. This may help to determine if the genes were horizontally transferred and, if so, between which species. The discovery of fumonisin production in three distantly related fungal genera shows that the production of fumonisins may be more widespread in the fungal kingdom than previously believed.

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Supplementary Table 1: Media composition

Media	Content
CYA	Agar (15 g/l), Yeast extract (5 g/l) (Biokar Diagnostics), Czapek Dox Broth [Saccharose (30 g/l), NaNO ₃ (3 g/l), K ₂ PO ₄ (1 g/l), TMS (1 ml/l), MgSO ₄ (50 mg/l), KCl (50 mg/l), FeSO ₄ (0.01mg/l)] (DIFCO)
CYAS	NaCl (5 g/l), Agar (15 g/l), Yeast extract (5 g/l) (Biokar Diagnostics), Czapek Dox Broth [Saccharose (30 g/l), NaNO ₃ (3 g/l), K ₂ PO ₄ (1 g/l), TMS (1 ml/l), MgSO ₄ (50 mg/l), KCl (50 mg/l), FeSO ₄ (0.01 mg/l)] (DIFCO)
DG18	Glycerol (180 g/l), TMS (0.8 ml/l), Chloramphenicol (0.1 g/l), Dichloran-Glycerol agar base [Glucose (10 g/l), Bacto-Peptone (5 g/l), Na ₂ H ₂ PO ₄ (1 g/l), MgSO ₄ (0.5 g/l), Dichloran (0.2 mg/l)] (OXOID)
DRYES	Sucrose (150 g/l), Yeast extract (20 g/l) (Biokar Diagnostics), Agar (20 g/l), TMS (1 ml/l), MgSO ₄ (0.5 g/l), Chloramphenicol (0.1 g/l), Rose Bengal (5 mg/l), Dichloran (0.2 mg/l)
MEA	Malt extract (20 g/l) (DIFCO), Glucose (20 g/l), Agar (20 g/l), Peptone (1 g/l), TMS (1 ml/l)
OAT	Oat meal (30 g/l), Agar (15 g/l), TMS (1 ml/l)
PCA	Potatoes (10 g/l), Carrots (10 g/l), TMS (0.5 ml/l), Agar (10 g/l)
PDA	Dextrose (20 g/l), Agar (15 g/l), Potato starch (4 g/l), TMS (1 ml/l)
V8	Campbell's V8 Juice (175 g/l), CaCO ₃ (3 g/l)
YES	Sucrose (150 g/l), Yeast extract (20 g/l), Agar (20 g/l), TMS (1 ml/l), MgSO ₄ (0.5 g/l),

TMS: Trace metals solution (Stock solution, ZnSO₄·7H₂O (10 g/l), CuSO₄·5H₂O (5 g/l))

Paper 9

”Review of secondary metabolites and mycotoxins from the
Aspergillus niger group”

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Review of secondary metabolites and mycotoxins from the *Aspergillus niger* group

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Abstract Filamentous fungi in the *Aspergillus* section *Nigri* (the black aspergilli) represent some of the most widespread food and feed contaminants known but they are also some of the most important workhorses used by the biotechnological industry. The *Nigri* section consists of six commonly found species (excluding *A. aculeatus* and its close relatives) from which currently 145 different secondary metabolites have been isolated and/or detected. From a human and animal safety point of view, the mycotoxins ochratoxin A (from *A. carbonarius* and less frequently *A. niger*) and fumonisin B₂ (from *A. niger*) are currently the most problematic compounds. Especially in foods and feeds such as coffee, nuts, dried fruits, and grape-based products where fumonisin-producing fusaria are not a problem, fumonisins pose a risk. Moreover, compounds such as malformins, naphtho- γ -pyrones, and bicoumarins (kotanins) call for monitoring in food, feed, and biotechnology products as well as for a better toxicological evaluation, since they are often produced in large amounts by the black aspergilli. For chemical differentiation/identification of the less toxic species the diketopiperazine asperazine can be used as a positive marker since it is consistently produced by *A. tubingensis* (177 of 177 strains tested) and *A. acidus* (47 of 47 strains tested) but never by *A. niger* (140 strains tested). Naphtho- γ -pyrones are the compounds produced in the highest quantities and are produced by all six common species in the group (*A. niger* 134 of 140; *A. tubingensis*

169 of 177; *A. acidus* 44 of 47; *A. carbonarius* 40 of 40, *A. brasiliensis* 18 of 18; and *A. ibericus* three of three).

Keywords Metabolomics · Fumonisin · Ochratoxin · Liquid chromatography–mass spectrometry · Polyketide synthase · Polyketide

Introduction

The black aspergilli are some of the most important mycotoxigenic food and feed contaminants, especially in postharvest decay of fresh and dried fruits and certain vegetables, nuts, beans, and cereals [1, 2]. This is due to their fast growth, pH tolerance, and high abundance in many environments.

For the analytical chemist, issues such as fungal taxonomy and correct identification may seem of low relevance, but in fact biosystematics is a vital part of mycotoxin research and food safety. Since the profile of mycotoxins and other secondary metabolites is species-specific [3–5], correct identification at the species level provides the key for planning the analytical determination of all relevant compounds.

The *Aspergillus niger* group (the black aspergilli, *Aspergillus* subgenus *Circumdati* section *Nigri*) comprises 18 species, of which *A. niger*, *A. tubingensis*, *A. brasiliensis*, *A. acidus*, *A. carbonarius*, and *A. ibericus* are common, whereas the remaining species are rare and found mainly in tropical regions (Table 1). A cladification of *Aspergillus* section *Nigri* using the β -tubulin and calmodulin genes showed that three clades could be distinguished [5]: the *A. niger* clade, a clade consisting of the two rare species *A. homomorphus* and *A. ellipticus*, and the clade of uniseriate black aspergilli (*A. aculeatinus*, *A. aculeatus*, *A. japonicus*,

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Table 1 Species in the *Aspergillus niger* group

Species	Rare	Species	Rare
<i>A. acidus</i> ^a		<i>A. ibericus</i>	
<i>A. brasiliensis</i> ^a		<i>A. niger</i>	
<i>A. carbonarius</i>		<i>A. piperis</i>	Yes
<i>A. costaricensis</i>	Yes	<i>A. scleroticarbonarius</i>	Yes
<i>A. ellipticus</i>	Yes	<i>A. sclerotioniger</i>	Yes
<i>A. heteromorphus</i>	Yes	<i>A. turingensis</i> ^a	
<i>A. homomorphus</i>	Yes	<i>A. vadensis</i>	Yes
Uniseriate black aspergilli			
<i>A. aculeatinus</i>		<i>A. japonicus</i>	
<i>A. uvarum</i>		<i>A. aculeatus</i>	

Species whose names are in **bold** have only been found in the tropics as yet. Important synonyms of *A. niger* are *A. awamori*, *A. phoenicis*, *A. kawachii*, *A. saitoi*, *A. usamii*, *A. foetidus*, *A. citricus*, and *A. ficuum*.

^a Regularly confused with *A. niger*

and *A. uvarum*), the members of which differ significantly from the remaining black aspergilli regarding their morphology, physiological behavior, and secondary metabolite production (e.g., producers of neoxaline, asperparalines, secalonc acids, asperamide, and aculeasins) [5], and this third clade has therefore not been included in this review. The identity and metabolite production of the uniseriate black aspergilli is usually not confused with the identity and metabolite production of *A. niger* and other biseriate black aspergilli.

A. niger and *A. turingensis* are probably the most common of the black aspergilli; however, in many studies describing secondary metabolites from these aspergilli, the producing organism has been identified as a black *Aspergillus* and then in many cases incorrectly named *A. niger*. A wrong identification may be further complicated by insufficient molecular identification based on sequencing of ribosomal DNA with low resolution [5, 6]. A polyphasic approach where many different types of characters (microscopy, metabolite profiling, molecular methods) are used is recommended for the identification of these aspergilli. Certain molecular methods have proven

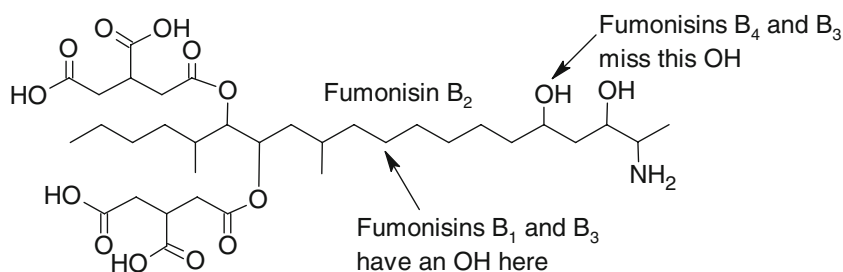
quite successful, including restriction fragment length polymorphism and β -tubulin, or calmodulin sequencing (reviewed in [5]).

Until recently, the main mycotoxin from the black aspergilli was considered to be ochratoxin A (OTA), produced in variable amounts within certain species of the group. *A. carbonarius* consistently produces large amounts of OTA, whereas only 6–10% of members of the *A. niger* group produce OTA and in 10–1000 lower amounts [7–9]. The third species producing OTA in section *Nigri* is *A. sclerotioniger*, but this species has only been found once in coffee. However, *A. niger* clearly has the widest distribution and has been reported to grow and damage a much larger number of crops and foods worldwide, including corn, peanuts, raisins, onions, mango, apples, and dried meat products [2]. This combined with the recent discovery that *A. niger* can also produce fumonisin B₂ (FB₂) and fumonisin B₄ (FB₄) [10, 11] (Fig. 1) necessitates the addition of fumonisins in a number of food and feed screening programs.

A. niger exhibits a remarkably versatile metabolism, which has made it one of the most important production organisms used for industrial fermentations [12, 13]. Since 1923, *A. niger* has been exploited commercially for its production of citric acid, mostly for use in food, cosmetics, and pharmaceutical preparations [14]. In addition, the fungus has been a rich source of industrial enzymes such as α -amylases, cellulases, and pectinases for use in the food industries since the 1960s [12]. *A. niger* possesses posttranslational mechanisms capable of correctly processing proteins that are difficult to express in traditional host organisms. As a result, it is widely used as a cell factory for heterologous expression of proteins [14].

A. niger has been considered to be nontoxic under industrial conditions [12], and thus to be considered a safe production organism. As a result, quite a number of *A. niger* fermentations have been granted the generally regarded as safe (GRAS) status by the US Food and Drug Administration [12]. However, the potential presence of both OTA and fumonisins in *A. niger* emphasizes the need to adjust and/or reconsider the screening procedures for simultaneous targeting of multiple classes of mycotoxins.

Fig. 1 Structure of fumonisin B₂ and the difference from the structures of fumonisin B₁, fumonisin B₃, and fumonisin B₄



In this review, we focus on the important secondary metabolites produced by members of the *A. niger* group relevant to the food, feed, and biotechnology industries. We have critically scrutinized the existing literature for reports of secondary metabolites claimed to be produced by *A. niger*, however often just found in a single strain or on a single occasion. In addition, we present analytical results from 25 years of metabolite profiling at the Center for Microbial Biotechnology of the black aspergilli. The strains investigated come from a large in-house collection (IBT collection, author address) and other fungal collections. The results are based on continuously obtained data from liquid chromatography (LC) with diode-array detection (DAD), 6 years of LC-DAD analyses combined with high-resolution time-of-flight (TOF) mass-spectrometric detection, and 1 year of screening with LC and tandem mass-spectrometric detection.

The overall aim is to provide an overview of the large numbers of compounds produced within this important group of filamentous fungi and to identify potential difficulties and pitfalls in the biochemical analysis of these compounds. Therefore, we discuss the individual compound classes, their biological significance, and natural abundance together with their spectroscopic and chromatographic properties relevant for their analytical determination.

Methodologies applied at the center for microbial biotechnology

Cultivation and extraction

Data obtained from analysis of *A. niger* group strains in our institution during the last 25 years were compiled. Cultures were identified on the basis of their morphology, metabolite profile, and partial sequencing (β -tubulin and calmodulin) as described elsewhere [15]. The strains were grown and extracted by one of the following three methods:

1. Combined chloroform–methanol (2:1, v/v) and acetone–ethyl acetate (1:1, v/v) extracts from cultures on yeast extract sucrose (YES) agar, Sigma YES agar, oatmeal agar, and potato sucrose agar (three plates each), as described by Frisvad and Thrane [16]. These were made in the years 1983–1995, and comprise approximately 150 fungal isolates.
2. Extracts made by the microextraction procedure of Smedsgaard [17], where approximately 0.6-cm² plugs of culture from YES agar, Czapek yeast autolysate agar (CYA), CYA with 5% salt (CYAS), oatmeal agar, or malt extract agar [4] were extracted with methanol–dichloromethane–ethyl acetate (1:2:3 v/v) and 1% formic acid. These comprise some 200 fungal isolates and 200 extracts made in the years 1996–2009.

3. Extracts for fumonisin analysis made by the microextraction procedure mentioned above, but extracted using 75% methanol [10] in the years 2007–2009. Here mainly CYAS and partly YES agar cultures were extracted, which add up to a further 100 cultures and 200 extracts.

Metabolite analysis by LC-DAD

The large extracts from the years 1983–1995 were analyzed by LC [acidic, 15–100% acetonitrile (CH₃CN) gradient, 40 min, Nucleosil C₁₈ column] with DAD (200–600 nm) as described by Frisvad and Thrane [16]. Data were available as printed reports with chromatographic traces (210 and 280 nm) and UV/vis spectra (200–600 nm).

Microextracts from the years 1995–2003 were analyzed by LC-DAD with parallel fluorescence detection (FLD) under similar chromatographic conditions using a Nucleosil BDB C₁₈ column [17] with the FLD set at 230→333 and 230→450 nm.

Microextracts from the years 2003–2009 were analyzed by LC-DAD-FLD as mentioned above using a Luna C₁₈ II column (15% CH₃CN to 100% CH₃CN in 20 min) [18].

Metabolite analysis by LC-DAD–TOF mass spectrometry

These analyses were performed using a LC system coupled to an orthogonal TOF mass spectrometer (Micromass LCT oaTOF) equipped with an electrospray source [19]. Two different gradients were used: (1) Luna II C₁₈ 15% CH₃CN to 100% CH₃CN in 20 min, 20 mM formic acid in both solvents [20]; (2) as described before but with the gradient starting at 30% CH₃CN and going to 60% CH₃CN in 5 min and then to 100% in 1 min.

About 300 extracts were analyzed by LC-DAD–TOF mass spectrometry (TOFMS) in the years 2003–2009; of these, most have also been analyzed by LC-DAD-FLD.

Metabolite analysis by LC–tandem mass spectrometry

This was performed using a Quattro Ultima triple quadrupole mass spectrometer equipped with an electrospray ionization (ESI) source and operated in multiple reaction mode (MRM) [11]. Separation was done on a Gemini C₆-phenyl column (acidic, 20% CH₃CN to 55% CH₃CN in 6 min, then to 100% in 30 s). MRM transitions for OTA, fumonisin B₁ (FB₁), FB₂, fumonisin B₃ (FB₃), and FB₄ are described elsewhere [11, 21]. Malformin A₂ was detected using m/z 516→304 at 30 V and m/z 516→417 at 20 V; malformins C and A were detected using m/z 530→372 at 25 V and m/z 530→417 at 20 V; ochratoxin β was detected using m/z 223→103 at 36 V and m/z 223→159 at 36 V; ochratoxin α was detected using m/z 257→193 at 36 V and

m/z 257→221 at 33 V; and ochratoxin B was detected using m/z 370→205 at 36 V and m/z 370→324 at 36 V. About 350 extracts were analyzed by LC–tandem mass spectrometry (MS/MS) in the years 2008–2009; many of these have also been analyzed by LC-TOFMS and LC-DAD-FLD.

Reference standards

For the LC-DAD-FLD analyses, OTA, ochratoxin B (Sigma-Aldrich), ochratoxin α , kojic acid, nigragillin, and malformins A, A₂, and C (gifts) were coanalyzed. For LC–mass spectrometry (MS), FB₁, FB₂, and FB₃ (Biopure, Tulln, Austria) and AAL toxins TA₁ and TB₁ (Sigma-Aldrich) were also included. Tensidols A and B and aurasperones A, B, and E were purified and validated by NMR [22].

Data analysis

LC-DAD and LC-DAD-FLD data were analyzed manually by comparing retention times, and UV spectra for the detected peaks as well as the retention times of the fluorescence peaks. These data were matched with LC-DAD-TOFMS data for tentative identification of the peaks by searching Antibase (Wiley) for compounds with the same accurate mass (± 0.01 -Da tolerance) and similar UV properties and estimated retention time (based on theoretical $\log D$ values) [19]. Compounds that we were able to tentatively identify in extracts from black aspergilli were aurasperones C–G, pyrophene, funalenone, fonsecinones B and C, fonsecin (TMC-256B1B), TMC-256A2, rubrofusarin B, fonsecin B, tensuic acid A, asperazine, pyranonigrins A–D, and FB₄. The identity of FB₄ was further validated by showing that its accurate mass, retention time and tandem spectra from 25–50 eV were identical to those of the presumed FB₄ in culture extracts from *Fusarium verticillioides*.

Analysis of biological significant metabolites

The black aspergilli can produce a diverse range of mainly polyketide-derived secondary metabolites along with non-ribosomal peptides and a number of compounds of mixed biosynthetic origin. Annotation of the fully sequenced genome of *A. niger* showed the presence of an impressive 34 polyketide synthase (PKS)-encoding genes as well as 17 nonribosomal peptide synthase (NRPS) genes and seven PKS-NRPS hybrids, accentuating the biosynthetic potential and versatility of this species [13]. So far, a total of 145 metabolites (Table 2) have been isolated and their structures elucidated from the biseriate black aspergilli.

Ochratoxins

Historically, OTA and its precursors (ochratoxins B, β , and α) have been the most significant mycotoxins produced within the *A. niger* complex. In extracts from approximately 400 agar cultures, OTA was always detected in amounts at least 10 times higher than the precursors using LC-FLD, LC-TOF analysis, and LC-MS/MS, and in many cases only OTA was detected.

It has been claimed that *A. tubingensis* can produce OTA [23, 24], presumably owing to unspecific chemical analyses. It was not possible to confirm this from the same cultures in our laboratory by LC-FLD, LC-TOF analysis, nor LC-MS/MS in culture extracts from up to ten different media. Other researchers have also doubted OTA production by *A. tubingensis* [3, 25].

Owing to the array of adverse toxicological effects on animals and humans, OTA is one of the most studied mycotoxins. The very low regulatory levels in food and feed mean that numerous highly sensitive analytical methods have been developed [26–28] and these have been reviewed several times [29, 30]. Thus, OTA will not be in the major scope of this paper since many of the commodities where *A. niger* is a problem have already been covered [29, 31–35]. However, it must be emphasized that we have found that direct analysis of OTA from *A. niger* using LC-DAD is not possible in crude extracts owing to coeluting malformins and naphtho- γ -pyrones (NGPs) (see later), of which the NGPs are present in more than 100 times higher quantities than OTA. This is illustrated in Fig. 2 showing LC-DAD-TOFMS chromatogram with the OTA peak hidden underneath the peaks of rubrofusarin B and malformins. Furthermore, with use of our LC-DAD-FLD method, four to five peaks of closely eluted substances are detected from many black *Aspergillus* extracts (230→333 nm, acidic conditions). This indicates a risk that some of these compounds may be coeluted with OTA even under other slightly different conditions. Thus, careful control of chromatographic conditions is necessary. Careful control of potential carryover between samples in the autosampler should be done if high-level OTA-producing *A. carbonarius* strains are analyzed in sequence.

For OTA analysis, our LC-TOFMS was approximately twofold less sensitive than FLD. Here sensitivity could be enhanced tenfold by shifting to alkaline conditions [32]; however, other compounds would suffer from a pH change (e.g., by postcolumn addition of NH₄OH solution) and comparison with old data will not be possible. Our LC-MS/MS instrument is about 20–50 times more sensitive than acidic FLD and has until now not yielded any interfering peaks on any of the two MRM transitions. It should be noted that the latest LC-TOFMS and Orbitrap instruments will, according to the manufactures, provide 10 times

Table 2 Secondary metabolites and mycotoxins from *A. niger* and related species

Component	Species	Elementary composition	UV/VIS	Monoisotopic mass (Da)
Glyoxylic acid [86, 87]		C ₂ H ₂ O ₃		74.0004
Glycolic acid [86]		C ₂ H ₄ O		76.0155
Oxalic acid [88]		C ₂ H ₂ O ₄		89.9947
Hydroxypyruvic acid [89]		C ₃ H ₄ O ₄		104.0104
(+)-Parasorbic acid (Antibase2008)		C ₆ H ₈ O ₂		112.0519
Sorbic acid (Antibase2008)		C ₆ H ₈ O ₂		112.0519
Fumaric acid [90]		C ₄ H ₄ O ₄		116.0104
2-Phenylethanol [91]		C ₈ H ₁₀ O		122.0726
Glutaric acid [86]		C ₅ H ₈ O ₄		132.0423
Asperyllone [92]		C ₂₀ H ₂₂ O		278.1671
Phenylacetic acid [91]		C ₈ H ₈ O ₃		136.0519
Phenoxyacetic acid [91]		C ₈ H ₈ O ₃		152.0468
<i>p</i> -Methoxyphenylacetic acid [91]		C ₉ H ₁₀ O ₃		166.0624
4-Hydroxymandelic acid [88]		C ₈ H ₈ O ₄		168.0417
Dehydrocarolic acid [93]	<i>A. brasiliensis</i>	C ₉ H ₈ O ₄		180.0417
Cyclo-L-Ala-L-Leu [94]		C ₉ H ₁₆ N ₂ O ₂		184.1206
Citric acid [95]	<i>Nigri</i> ^a	C ₆ H ₈ O ₇		192.0265
D-Galactonic acid		C ₆ H ₁₂ O ₇		196.0583
7-Hydroxy-4-methoxy-5-methylcoumarin [96, 97]	<i>A. niger</i> (reported as <i>Cladosporium herbarum</i>)	C ₁₁ H ₁₀ O ₄		206.0574
Hexylitaconic acid [98, 99]	<i>A. niger</i>	C ₁₁ H ₁₈ O ₄		214.1200
3-Methyl-8-hydroxy-4-decanoate [91]	<i>A. niger</i>	C ₁₂ H ₂₂ O ₃		214.1564
Ochratoxin β	<i>A. niger</i> , <i>A. carbonarius</i> , <i>A. sclerotiumniger</i>	C ₁₁ H ₁₀ O ₅		222.0523
Nigragillin [93]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₁₃ H ₂₂ ON ₂	262 (26,200)	222.1739
Pyranonigrin A (pyranopyrrol A) [57, 75]	<i>Nigri</i>	C ₁₀ H ₉ NO ₅	210, 252, 316	223.0481
Aspernigrin A [57, 96]	<i>A. niger</i>	C ₁₃ H ₁₂ N ₂ O ₂		228.0893
Carbonarone A [83]	<i>A. carbonarius</i>	C ₁₃ H ₁₁ NO ₃		229.0733
Carbonarone B [83]	<i>A. carbonarius</i>	C ₁₃ H ₁₁ NO ₃		229.0733
Carbonarone A [83]	<i>A. carbonarius</i>	C ₁₃ H ₁₁ NO ₃	279 (100%), 317 (29%), 401 (25%)	229.0739
Carbonarone B [83]	<i>A. carbonarius</i>	C ₁₃ H ₁₁ NO ₃	340 (100%)	229.0739
Tensidol A [100]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₁₃ H ₁₁ NO ₃	206 (100%), 250 (30%)	229.0739
2-Methylene-3-hexylbutanedioic acid [98]	<i>A. niger</i>	C ₁₁ H ₁₈ O ₅		230.1149
Pyranonigrin D (pyranopyrrol D) [57]	<i>Nigri</i>	C ₁₁ H ₉ NO ₅		235.0481
2-Carboxymethyl-3-hexylmaleic acid anhydride [98, 101]	<i>A. niger</i>	C ₁₂ H ₁₆ O ₅	253	240.0992
Tensyuic acid [102]	<i>A. niger</i>	C ₁₁ H ₁₆ O ₆	203 (6,000), 205 (5,600)	244.0947
Tensyuic acid F [102]	<i>A. niger</i>	C ₁₁ H ₁₆ O ₆	199 (11,000)	244.0947
Pyranonigrin B (pyranopyrrol B) [57]	<i>Nigri</i>	C ₁₁ H ₁₁ NO ₆		253.0586
Pyranonigrin C (pyranopyrrol C) [57]	<i>Nigri</i>	C ₁₁ H ₁₁ NO ₆		253.0586
Ochratoxin α	<i>A. carbonarius</i> , <i>A. niger</i> , <i>A. sclerotiumniger</i>	C ₁₁ H ₉ ClO ₅		256.0133
Tensyuic acid B [102]	<i>A. niger</i>	C ₁₂ H ₁₈ O ₆	204 (8,900), 207 (7,700)	258.1103
Nigerazine A [71]	<i>A. niger</i> ?	C ₁₆ H ₂₂ N ₂ O		258.1727
Nigerazine B [71]	<i>A. niger</i> ?	C ₁₆ H ₂₂ N ₂ O	280 (22,400)	258.1732
Nigerloxin [79]	<i>A. niger</i> ?	C ₁₃ H ₁₅ NO ₅		265.0945
TAFU-567 [103]	<i>A. acidus</i> ?	C ₁₃ H ₁₄ O ₆		266.0785
Antafumicin A [104]	<i>A. acidus</i>	C ₁₃ H ₁₄ O ₆	217(21,000), 279(15,500), 317(8,720)	266.0786

Table 2 (continued)

Component	Species	Elementary composition	UV/VIS	Monoisotopic mass (Da)
Antafumicin B [104]	<i>A. acidus</i>	C ₁₃ H ₁₄ O ₆	217(21,000), 279(15,500), 317(8,720)	266.0786
TMC-256A1 [105]	<i>A. niger</i>	C ₁₅ H ₁₂ O ₅	236 (100%), 276 (68%), 368 (12%), 482 (1%)	272.0679
TMC-256C1 [105]	<i>A. niger</i>	C ₁₅ H ₁₂ O ₅		272.0679
Rubrofusarin(heminigerone) [44, 106]	<i>Nigri</i>	C ₁₅ H ₁₂ O ₅	(225, 28,200) (253sh, 19,600) (278, 49,700) (328, 3,400) (415, 5,200)	272.0679
Tensyucic acid C [102]	<i>A. niger</i>	C ₁₃ H ₂₀ O ₆	201 (7,700), 206 (4,500)	272.1260
Tensyucic acid D [102]	<i>A. niger</i>	C ₁₃ H ₂₀ O ₆	202 (17,000), 205 (7,700)	272.1260
Asperribol [64, 107]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₀ H ₂₂ O	242(9,800), 293(11,200), 400(63,200)	278.1672
Asperic acid [65]	<i>A. niger</i>	C ₁₆ H ₂₈ O ₄	225, 280	284.1988
Flavasperone [108, 109]	<i>Nigri</i>	C ₁₆ H ₁₄ O ₅	225(28,184), 254(47,863), 406(5,495)	286.0841
Tensyucic acid E [102]	<i>A. niger</i>	C ₁₄ H ₂₂ O ₆	198 (13,000)	286.1416
Pyrophen [65]	<i>A. tubingensis</i> , <i>A. niger</i>	C ₁₆ H ₁₇ NO ₄		287.1158
Funalenone [55]	<i>A. niger</i> , <i>A. tubingensis</i> , <i>A. brasiliensis</i>	C ₁₅ H ₁₂ O ₆		288.0634
Fonsecin (TMC-256B1) [110]	<i>Nigri</i>	C ₁₅ H ₁₄ O ₆		290.0790
Nigerasperone A [111]	<i>A. niger?</i>	C ₁₆ H ₁₄ O ₆	210sh (55%), 226 (74%), 255sh (56%), 276 (100%), 325 (14%), 400 (22%)	302.0785
Xanthoherquein [55]	<i>A. sclerotiumniger?</i>	C ₁₅ H ₁₂ O ₇		304.0578
Fonsecin B (TMC-256B2) [105, 110, 112]	<i>Nigri</i>	C ₁₆ H ₁₆ O ₆	231 (80%), 275 (100%), 328 (26%), 406 (21%)	304.0941
Fonsecin monomethyl ether [44, 110, 112]	<i>Nigri</i>	C ₁₆ H ₁₆ O ₆	232(28,500), 277(40,500), 317(9,100), 330(10,000), 395(8,400)	304.0947
Asnipyrone B [113]	<i>A. niger?</i>	C ₂₀ H ₂₀ O ₃	235(16,600), 265(15,490), 376(33,885)	308.1413
Carbonarin E [114]	<i>A. carbonarius</i>	C ₁₈ H ₁₇ NO ₄		311.1158
Asperenone [107]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₂ H ₂₄ O ₃	242(13,800), 300(9,200), 414–416(94,200)	312.1726
4,9-Dihydroxypyrone-3,10-quinone [88]	<i>Nigri?</i>	C ₂₀ H ₁₀ O ₄		314.0573
Asnipyrone A [113]	<i>A. niger</i>	C ₂₁ H ₂₂ O ₃	240(18,600), 290(19,055), 378(31,625)	322.1570
Atromentin [75]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₁₈ H ₁₂ O ₆		324.0628
Cycloleucomelon [57]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₁₈ H ₁₀ O ₇		338.0421
Tensidol B [100]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₁₈ H ₁₇ NO ₆	206 (100%),	343.1050
Nafuredin [115]	<i>A. niger?</i>	C ₂₂ H ₃₂ O ₄		360.2295
22-Deacetyl-anuthone A [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₂ H ₃₂ O ₄	238(16,220), 302(480)	360.2302
Aspergillin [54]	<i>A. Nigri</i>	C ₁₉ H ₁₂ O ₈		368.0527
2,3,4,8,9,10-Hexahydroxy-5H,6H,7H-benzopyrene-1,11-dione [88]	<i>Nigri</i>	C ₁₉ H ₁₂ O ₈		368.0527
1-Hydroxy-anuthone A [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₄ H ₃₂ O ₅	234(8,510), 290(220)	400.2251
1-Hydroxy-anuthone C [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₄ H ₃₂ O ₅	232(4,680), 282(468)	400.2251
Yanuthone B [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₄ H ₃₂ O ₅	244(4,470), 290(1,950), 338(830)	400.2251
Tubingensis A [116]	<i>A. tubingensis</i> ATCC 76608 ^b	C ₂₈ H ₃₅ NO		401.2713
Tubingensis B [117]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₅ NO	218(17,200), 237(25,500), 260(10,100), 299(10,100), 325(2,200), 338(6,700)	401.2720
Yanuthone A [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₄ H ₃₄ O ₅	232(7,950), 310(500)	402.2407
Yanuthone C [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₄ H ₃₄ O ₅	232(13,180), 280(2,040), 322(1,070)	402.2407
Ochratoxin A [118]	<i>A. niger</i> , <i>A. carbonarius</i> , <i>A. sclerotiumniger</i>	C ₂₀ H ₁₈ ClNO ₆	205(94%), 215(98%), 218(21%), 283(2%), 332(17%)	403.0823
Dihydrotubingensis B [119]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₇ NO		403.2870
Dihydrotubingensis A [119]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₇ NO		403.2870
10,23-Dihydro-24,25-dehydroaflavinine [120]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₉ NO	226(31,600), 284(4,400), 291(4,100)	405.3032
Aflavinine [120]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₉ NO	225(18,700), 283(3,010), 291(2,710)	405.3032
Bicoumanigrin (JH 0508 O'; bicumarine) [57]	<i>A. niger</i>	C ₂₂ H ₁₈ O ₈		410.0996

Table 2 (continued)

Component	Species	Elementary composition	UV/VIS	Monoisotopic mass (Da)
Orlandin [56]	<i>A. niger</i>	C ₂₂ H ₁₈ O ₈	311(29,500), 321(25,120)	410.1002
BMS-192548 [52, 121]	<i>A. niger</i> ?	C ₂₁ H ₁₈ O ₉	280(4,440), 320(35,548), 414(38,822)	414.0950
Nygerone A [76]	<i>A. niger</i>	C ₂₄ H ₂₂ N ₂ O ₅		418.1529
10,23-Dihydro-24,25-dehydro-21-oxo-aflavinine [120]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₇ NO ₂	224(18,200), 267(4,200), 273(1,400), 283(2,100), 291(1,750)	419.2824
14-epi-14-hydroxy-10,23-dihydro-24,25-dehydroaflavinine [122]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₉ NO ₂		421.2975
14-epi-14-hydroxy-10,23-dihydro-24,25-dehydro-aflavinine [120]	<i>A. tubingensis</i> ATCC 76608	C ₂₈ H ₃₉ NO ₂	226(16,400), 284(2,290), 292(2,100)	421.2980
Demethylkotanin [75]	<i>A. niger</i>	C ₂₃ H ₂₀ O ₈		424.1153
Aspernigerin [73]	<i>A. niger</i>	C ₂₆ H ₃₂ N ₄ O ₂		432.2525
Kotanin [56]	<i>A. niger</i>	C ₂₄ H ₂₂ O ₈	203(92%), 208(100), 237(35), 259(15%), 296(36%), 306(41%), 317(35%)	438.1309
Aspernigrin B [57, 96]	<i>A. niger</i>	C ₂₇ H ₂₄ N ₂ O ₅		456.1679
Yanuthone D [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₈ H ₃₈ O ₈	226(7,940), 254(3,715), 298(1,150)	502.2567
Malformin A ₂ [63, 67]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₂ H ₃₇ O ₅ N ₅ S ₂	End abs.	515.2239
Malformin B ₅ [63, 67]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₂ H ₃₇ O ₅ N ₅ S ₂	End abs.	515.2239
Malformin B ₂ [63, 67]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₂ H ₃₇ O ₅ N ₅ S ₂	End abs.	515.2239
Yanuthone E [77]	<i>A. niger</i> , <i>A. tubingensis</i>	C ₂₉ H ₄₂ O ₈	234(9,120), 280(1,000), 340(450)	518.2881
Malformin A ₁ [63]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₃ H ₃₉ O ₅ N ₅ S ₂	End abs.	529.2396
Malformin B _{1a} [63, 67]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₃ H ₃₉ O ₅ N ₅ S ₂	End abs.	529.2396
Malformin B _{1b} [63, 67]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₃ H ₃₉ O ₅ N ₅ S ₂	End abs.	529.2396
Malformin B ₅ [63, 67]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₃ H ₃₉ O ₅ N ₅ S ₂	End abs.	529.2396
Malformin B ₃ [63, 67]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₃ H ₃₉ O ₅ N ₅ S ₂	End abs.	529.2396
Malformin C [123]	<i>A. niger</i> , <i>A. brasiliensis</i> , <i>A. tubingensis</i>	C ₂₃ H ₃₉ O ₅ N ₅ S ₂	End abs.	529.2396
8'-O-Demethylnigerone [49]	<i>A. niger</i>	C ₃₁ H ₂₄ O ₁₀	284 (56,200), 365 (12,020), 400 (9,775)	556.1369
8'-O-Demethylisonigerone [49]	<i>A. niger</i>	C ₃₁ H ₂₄ O ₁₀	281 (33,885), 355 (26,915), 389 (7,760)	556.1369
Asperpyrone A [124]	<i>Nigri</i>	C ₃₁ H ₂₄ O ₁₀		556.1369
Dianhydroaurasperone C [125]	<i>Nigri</i>	C ₃₁ H ₂₄ O ₁₀	225(22,908), 255(26,915), 280(85,113), 325(3,715), 405(5,754)	556.1369
6'-O-Demethylnigerone [106]	<i>Nigri</i>	C ₃₁ H ₂₄ O ₁₀	226(39,810), 278(63,095), 408(10,000)	556.1369
Aurasperone D [44, 48, 102, 108, 109]	<i>Nigri</i>	C ₃₁ H ₂₄ O ₁₀	235–240(50,119), 280(51,286), 320–325(15,136), 380(7,080)	556.1369
Carbonarin C [114]	<i>A. carbonarius</i>	C ₃₃ H ₂₇ NO ₈		565.1737
Carbonarin D [114]	<i>A. carbonarius</i>	C ₃₃ H ₂₇ NO ₈		565.1737
Carbonarin H [114]	<i>A. carbonarius</i>	C ₃₃ H ₂₆ O ₉		566.1577
Isonigerone [49]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀	UV: 226 (65%), 275 (100%), 327 (17%), 407 (19%)	570.1526
Asperpyrone B [124]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀		570.1526
Asperpyrone C [124]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀		570.1526
Aurasperone A [47, 102, 108, 109]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀	225(51,300), 258(53,700), 280(100,000), 325(8,710), 406(12,880)	570.1526
Fonsecinone A [126]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀	228(45,700), 256(44,670), 278(67,600), 325(11,480), 398(7,080)	570.1526
Fonsecinone D [126]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀	227(26,900), 279(48,975), 316(8,128), 328(6,920), 403(6,310)	570.1526
Isoaurasperone A [44, 112]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀	255(33,884), 275(38,905), 385–390(3,467)	570.1626

Table 2 (continued)

Component	Species	Elementary composition	UV/VIS	Monoisotopic mass (Da)
Isonigerone [49, 106]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀	228(50,119), 248(56,234), 279(66,069), 390(9,550)	570.1626
Nigerone [106]	<i>Nigri</i>	C ₃₂ H ₂₆ O ₁₀	226(51,286), 278(83,186), 407(14,125)	570.1626
Nigerasperone C [111]	<i>Nigri</i>	C ₃₁ H ₂₆ O ₁₁	232 (50%), 282 (100%), 329 (15%), 403 (15%)	574.1470
Aurasperone F [48]	<i>Nigri</i>	C ₃₁ H ₂₆ O ₁₁	213, 281, 320, 334	574.1475
Carbonarin A [114]	<i>A. carbonarius</i>	C ₃₃ H ₂₆ O ₁₀		582.1526
Carbonarin B [114]	<i>A. carbonarius</i>	C ₃₃ H ₂₆ O ₁₀		582.1526
Diketopiperazine dimer [68]	<i>A. tubingensis</i>	C ₃₃ H ₃₈ N ₆ O ₄	240, 300	582.2949
Aurasperone G [48]	<i>Nigri</i>	C ₃₁ H ₂₄ O ₁₂	UV: 213, 281, 315, 332, 403	588.1268
Aurasperone E [44]	<i>Nigri</i>	C ₃₂ H ₂₈ O ₁₁	230(72,444), 282(147,910), 322(22,387), 330(20,418), 400(19,500)	588.1631
Fonsecinone B [126]	<i>Nigri</i>	C ₃₂ H ₂₈ O ₁₁	229(38,900), 255(41,700), 280(70,800), 320(14,125), 328(14,125), 403(9,120)	588.1631
Fonsecinone C [126]	<i>Nigri</i>	C ₃₂ H ₂₈ O ₁₁	234(51,285), 254(38,000), 279(60,255), 315(19,500), 327(14,800), 398(7,080)	588.1631
Aurasperone C [44, 47, 102, 108, 109]	<i>Nigri</i>	C ₃₁ H ₂₈ O ₁₂	236, 283.5, 323, 336, 412	592.1581
Carbonarin G [114]	<i>A. carbonarius</i>	C ₃₄ H ₂₈ O ₁₀		596.1682
10,10'-Bifonsecin B [49]	<i>Nigri</i>	C ₃₂ H ₃₀ O ₁₂	231 (84%), 276 (100%), 330 (26%), 411 (24%)	606.1732
Nigerasperone B [111]	<i>Nigri</i>	C ₃₂ H ₃₀ O ₁₂	236 (98%), 281 (100%), 316sh (32%), 380 (12%)	606.1732
Aurasperone B [44, 47, 48, 108, 109]	<i>Nigri</i>	C ₃₂ H ₃₀ O ₁₂	235(51,000), 282(87,700), 321(20,100), 334(20,000), 410(13,600)	606.1732
Carbonarin F [114]	<i>A. carbonarius</i>	C ₃₄ H ₂₈ O ₁₁		612.1632
Asperazine [64]	<i>A. tubingensis</i>	C ₄₀ H ₃₆ N ₆ O ₄	225, 275, 300	664.2798
Fumonisin B ₄ [11]	<i>A. niger</i>	C ₃₄ H ₅₉ NO ₁₃	Not UV active	689.3986
Fumonisin B ₂ [10]	<i>A. niger</i>	C ₃₄ H ₅₉ NO ₁₄	Not UV active	705.3930
Biotransformation products (not true metabolites)				
Differenol A (genistein, prunetol, sophoricol) [127]		C ₁₅ H ₁₀ O ₅		270.0528
3',4',5,7-Tetrahydroxy-8-methoxy isoflavone [128]		C ₁₆ H ₁₂ O ₇		316.0578
Pisolithin B [88]		C ₈ H ₈ O ₄		168.0417
8-Hydroxygenistein [128]		C ₁₅ H ₁₀ O ₆		286.0472
Flaviolin [129]		C ₁₀ H ₆ O ₅		206.0215
Orobole [127]		C ₁₅ H ₁₀ O ₆		286.0472
Iso-T-2 toxin (Antibase2008)		C ₂₄ H ₃₄ O ₉		466.2197

Extinctions coefficient

The identities of species marked with a *question mark* have not been confirmed.

^a Series *Nigri*: *A. niger*, *A. tubingensis*, *A. acidus*, *A. brasiliensis*, *A. carbonarius*, *A. sclerotivarbonarius*, *A. sclerotioniger*, *A. ibericus*, *A. vadensis*, *A. costaricaensis*, *A. piperis*

^b This isolate produces an abundance of sclerotia, but has not been unequivocally placed in *A. tubingensis* yet.

End abs. End absorption (UV max. <200nm)

higher mass resolution and 10–100 times better sensitivity than the LC-TOFMS instrument used here.

ESI of OTA mainly yields the protonated molecular ion [M+H]⁺ in positive mode and [M-H][−] in negative mode, with approximately 10 times higher signal in positive mode. With high in-source fragmentation settings, OTA yields several diagnostic ions for further confirmation. An unidentified fragment ion at *m/z* 358 can be observed along

with the sodium adduct [M+Na]⁺ and the ³⁷Cl [M+H]⁺ at *m/z* 406 [19]. Care should be taken when using nominal-mass LC-MS, since a very common contaminant from plasticware (presumably a phthalate) with a molecular mass of 386 Da is eluted very close to OTA and makes at predominant [M+NH₄]⁺ ion at *m/z* 404. OTA can be differentiated from the contaminant on the basis of the chlorine isotope pattern.

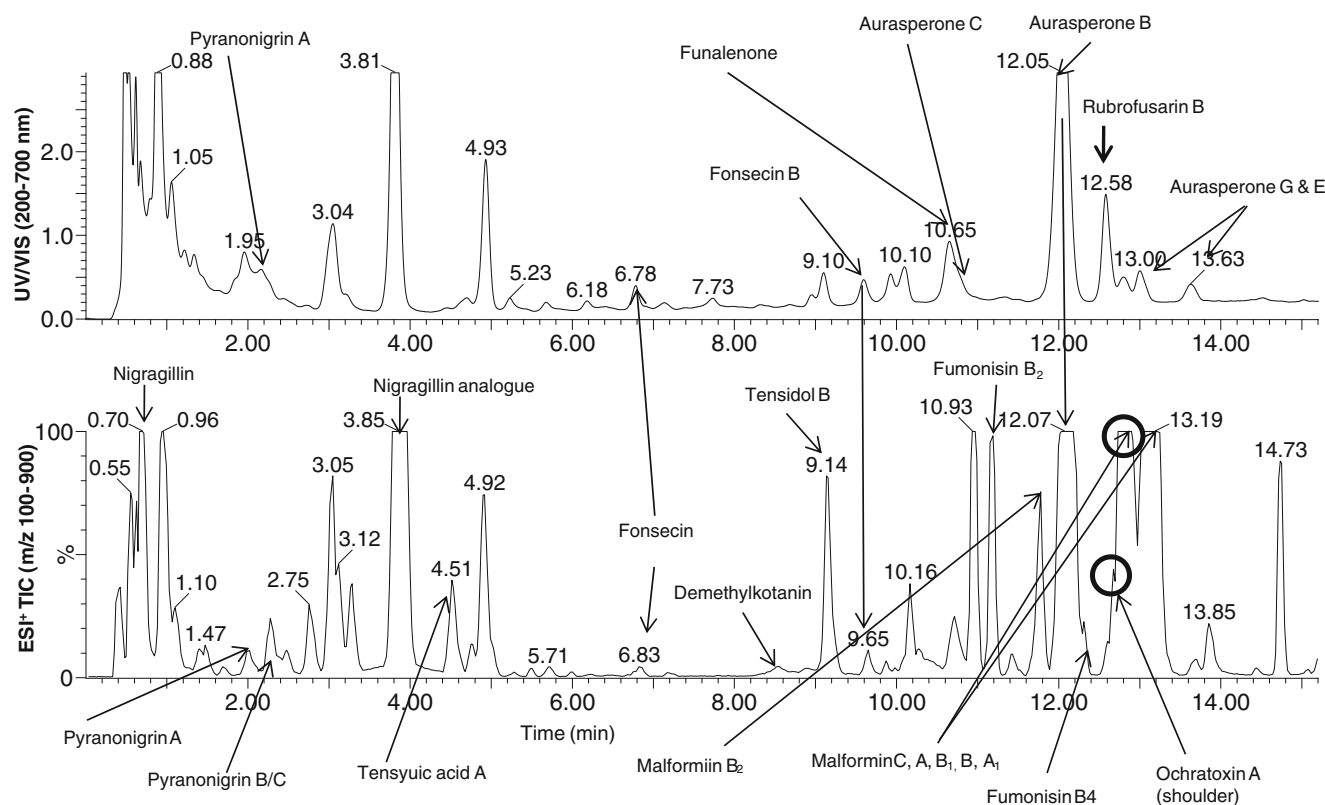


Fig. 2 *Aspergillus niger* extract from 8-day culture on yeast extract sucrose agar analyzed by liquid chromatography–diode-array detection–time-of flight mass spectrometry. The upper trace is the UV/vis (200–700 nm) chromatogram and the lower trace is the total ion

chromatogram (TIC) [positive electrospray ionization (ESI⁺), m/z 100–900]. Separation was done using a Luna C₁₈ II column with 15% CH₃CN to 100% CH₃CN in 20 min

Fumonisin

The fumonisins are a group of polyketide-derived mycotoxins, first discovered in 1988 from *F. verticillioides* [36]. This group of compounds is of great importance as they are suspected to be carcinogenic to humans, and are as a consequence regulated mainly in maize-based products. The fumonisins can be divided into four series A, B, C, and P [37], with the B series, mainly FB₁, FB₂, and FB₃ (Fig. 1), as the most abundant naturally occurring fumonisins [37, 38]. The surprising discovery of putative homologues to the *F. verticillioides* fumonisin gene cluster in two different *A. niger* genomes [13, 26] led to the subsequent documentation of actual FB₂ production in *A. niger* [10, 11]. Besides FB₂, FB₄ is also produced in lower amounts, about 10–25% of the amount of FB₂. From a current screening project on grapes and raisins it appears that fumonisins are produced by approximately 75% of all *A. niger* strains, whereas for coffee the figure was 76%, and they are thus much more common within *A. niger* than OTA production (6–10% [7, 8]).

Analysis of fumonisins produced by *Fusarium* spp. has previously been extensively reviewed [29, 37, 38]. In this review, FB₂ and FB₄ will therefore only be dealt with briefly in respect to the emerging data from analysis of

A. niger infected commodities for which new analytical methods needs to be developed. Owing to the lack of suitable chromophores or fluorophores, fumonisins cannot be detected directly using UV detection or FLD. This is probably why they were not detected in culture extracts of *A. niger* related species until recently. The fumonisins ionize excellently in positive ESI (ESI⁺) and also quite well in negative ESI (ESI[−]), yet about approximately tenfold lower than in ESI⁺ using our instruments. Bartok et al. [37] have listed comprehensive LC-MS/MS data on the fumonisins.

With the increasing sensitivity of LC-MS/MS, especially with triple quadrupole mass spectrometers, multimethods with no purification are emerging for low parts per billion levels of well-ionizing mycotoxins. An example is the screening method of Sulyok et al. [39, 40] which was used to detect fumonisins down to 3–17 µg/kg and OTA down to approximately 1 µg/kg. The method was used to detect very high levels of FB₂ (33 mg/kg) in dark bread, where the absence of FB₁ as well as a black infecting fungus on the bread together indicate that the contaminant was an *A. niger* rather than a *Fusarium* species. FB₂ and FB₄ probably produced by *A. niger* were also reported in [41] to have been detected by LC-TOFMS.

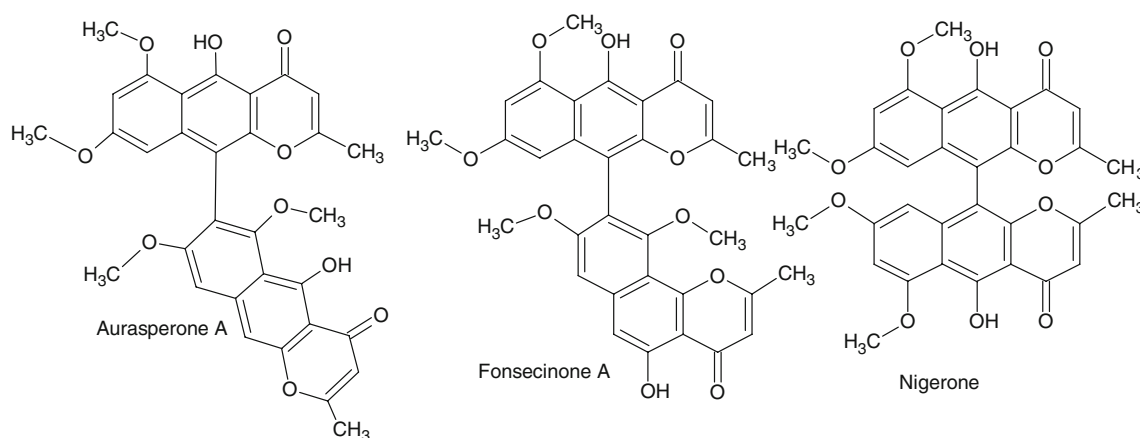


Fig. 3 Structures representing the three major groups of bis(naphtho- γ -pyrones) in the *A. niger* group

From CH_3CN extracts of strawberries, we were able to detect fumonisins with LC-MS/MS down to 10 $\mu\text{g/kg}$ and OTA down to 1 $\mu\text{g/kg}$, and on berries artificially infected with *A. niger* we found up to 25 mg/kg FB_2 and 2 mg/kg FB_4 , as well as up to 1 $\mu\text{g/g}$ OTA, whereas malformins were barely detected (K.F. Nielsen, unpublished results).

For analysis of green coffee, the EN 13585:2001 protocol (70% CH_3OH , strong anion exchange, SAX) for extraction and purification of fumonisins in maize could be directly adapted [11]. Furthermore, OTA which cooccurred in the extract was simultaneously retained and detected by LC-MS/MS [21]. Immunoaffinity purification of fumonisins was not a successful strategy for extracts from green and roasted coffee and resulted in viscous dark samples. A commercial ELISA kit (developed for maize) was also tested, but in green coffee it gave totally random results with many false positives and absolutely no correlation to LC-MS/MS results [11].

Since some *A. niger* strains produce very high amounts of citric acid and other small organic acids (up to 50 g/L) under some conditions, it has not always been possible to use SAX purification. Recoveries from SAX have been seen as low as 0% for OTA and fumonisins owing to competition with small organic acids. In consequence, we are currently testing strong cation exchange. Surprisingly, we have not found publications on cation-exchange purification of fumonisins, which is probably a reflection of the first fumonisin methods,

where derivatization with *O*-phthalaldehyde and subsequent LC-FLD was the preferred analytical strategy [29, 37, 38]. And since both *O*-phthalaldehyde and cation exchange target amines, these two methods are not orthogonal and thus not selective as a pair. However, with LC-MS as the detection principle, cation exchange should statistically be preferred since there are far fewer basic than acidic compounds in microorganisms.

Naphtho- γ -pyrones

Quantitatively, the NGPs represent the most abundant family of secondary metabolites in the *A. niger* group under all conditions observed (including chemostate cultures).

The biological effects of several NGPs have been investigated in various systems [14], and they have, for instance, been shown to be antibacterial, antifungal [42], antitumoral [43], and cytotoxic [42, 43]. Ghosal et al. [44] reported acute toxicity (interperitoneal) at the 10–50 mg/kg level; however, to the best of our knowledge no data on the bioavailability of these compounds exist. Thus, these compounds cannot currently be considered mycotoxins *sensu stricto*, since this requires toxicity via a natural route of exposure.

The NGP group of compounds comprises a series of aurasperones, Fonsecainones, and nigerones, as well as monomers such as flavasperone and rubrofusarin B (Figs. 3 and 4). Reanalysis of in-house LC-DAD and LC-MS data showed

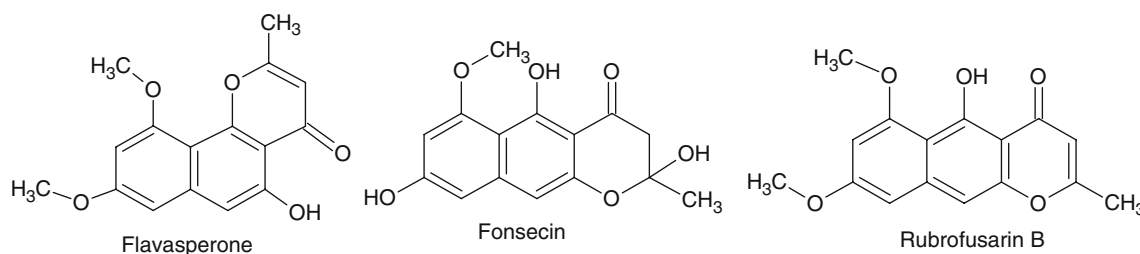


Fig. 4 Structures representing the three major groups of monomeric naphtho- γ -pyrones in the *A. niger* group

that all species within the “*A. niger* clade” are able to produce NGPs (*A. acidus*, *A. brasiliensis*, *A. carbonarius*, *A. costaricaensis*, *A. ibericus*, *A. niger*, *A. piperis*, *A. scleroticarbonarius*, *A. sclerotioniger*, *A. tubingensis*, and *A. vadensis* [5]), whereas species not producing NGPs include the “*A. ellipticus/A. heteromorphus* clade” and the “*A. aculeatus* clade” (comprising *A. homomorphus*, *A. aculeatinus*, *A. aculeatus*, *A. japonicus*, and *A. uvarum*). However, in losing the ability to produce black conidia, *A. laticoffeatus*, a light-brown-spored naturally occurring mutant of *A. niger*, has also lost the ability to produce NGPs, which is also the case for white-spored industrial strains we have analyzed. So there appears to be a link between black melanin production and NGP production in the *A. niger* group. Moreover, NGPs have been shown to be present in aerosolized spores [45].

Of 140 *A. niger* isolates we have analyzed by LC-DAD and LC-DAD-TOFMS, 134 (96%) were found to produce at least one NGP (usually aurasperone B in the highest amount). Comparable levels were seen for *A. tubingensis* (169 of 177, 95%), *A. ibericus* (three of three, 100%), and *A. acidus* (44 of 47, 94%) [46]. These data are thus conflicting with data reported by Bouras et al. [47], who only found NGPs in a low proportion of the strains investigated. This can be ascribed to differences in strain age, medium, and incubation conditions, since it has been our experience that the NGP levels are highest in fresh isolates and may decrease somewhat over time.

With LC-ESI-TOFMS, simple ionization patterns have been observed. In ESI⁺, [M+H]⁺ was the most abundant ion, and [M+Na]⁺ and [M+Na+CH₃CN]⁺ were the main adduct ions. In ESI[−], mainly [M−H][−] and low-abundance [M−2H+Na][−] ions was observed, the latter ion matching an acidic functionality (here a phenol group). Identical patterns were observed by Bouras et al. [47, 48], who also reported numerous specific fragment ions (aurasperones F and G) from MS/MS experiments (ion trap). These can also serve as potential MRM transitions for trace analysis.

The NGPs contain a fully conjugated system, giving rise to very characteristic UV/VIS spectra (Table 2, Fig. 5). However as seen in Table 2, and in the figures in Zhang et al. [49], some spectra are similar, and for the ones with the same elementary composition (e.g., ten different compounds with the formula C₃₂H₂₆O₁₀) it may even be difficult to separate them by their MS/MS spectra as they are mostly positional isomers which will probably fragment in a similar way. Thus, the retention time is the only way to differentiate NGPs if no reference standards are available or if no NMR validation can be done. An alternative is to compare the whole profiles with the one in Fig. 2 and the ones in Zhang et al. [49], since the order of elution of the NGPs should be the same if low-pH reversed-phase LC is used.

Since the NGPs constitute a large proportion of the total peak area and they absorb strongly in the 200–450-nm region, they often obscure the detection of compounds such as the malformins and ochratoxins. During preparative isolation of fumonisins, it has been observed that NGPs are not retained on SAX columns, but are retained on mixed-mode reversed-phase SAX columns (e.g., Oasis MAX), as secondary interactions with a polymeric backbone will cause strong retention of these weakly acidic phenols [50, 51]. These columns can then serve as a prefractionation step to selectively remove the NGPs.

Study of the occurrence of NGPs and other related compounds in food and feed samples is limited to the study of Ghosal et al. [44], who detected flavasperone, rubrofusarin, isoaurasperone, and aurasperones E, A, and D in artificially infected mangos at a total level of 60 mg/kg. NGPs were detected using thin-layer chromatography (280-nm absorption) after organic extraction and aurasperones A and D were found to be the predominant ones (approximately 33% each). Aurasperones (erroneously named tetracyclic compounds [52]) and orlandin were also found from *A. niger* infected building materials [53]. Since other metabolites originating

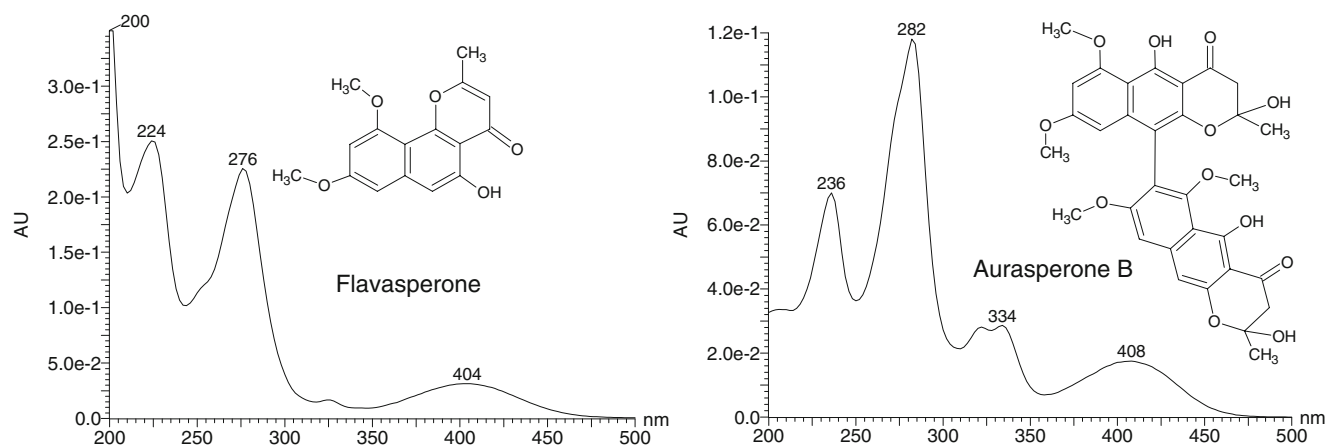
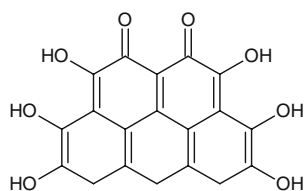


Fig. 5 UV spectra (pH 3.2) of monomeric (flasperone) and dimeric (aurasperone B) naphtho-γ-pyrones

Fig. 6 Structure of the green pigment aspergillin



from the *A. niger* group can be found in food (OTA and FB₂), NGPs are most likely to be naturally occurring in various food matrices, and hereby it would be recommendable to develop analytical protocols for detection of NGPs in such food types as well as investigating their bioavailability and subsequent toxicity.

Occasionally, we have been asked to analyze yellow samples from submerged fermentations with different, mainly industrial *A. niger* strains. With LC-DAD-TOFMS we have each time detected aurasperone B as the only secondary metabolite in the sample. In most cases, ESI[−] has been necessary for MS verification since foam suppressors [silicon and/or poly(ethylene glycol) oils] also suppress ESI⁺ signals.

In addition, a green pigment has been observed in *A. niger* fermentations. This is aspergillin (Fig. 6), a highly oxygenated polyketide with a hexahydroxyl pentacyclic quinoid structure [54]. Together with another kind of melanin made from monomers, the black pigment is made up from a complicated mixture of aspergillin and proteins [54].

Bicoumarins

The bicoumarins (Fig. 7) represent another prominent family of compounds from *A. niger*, consisting of a group of heterocyclic dimers derived from cinnamic acid lactone, further categorized on the basis of the type of connection between the coumarin moieties. Like the NGPs, the bicoumarins contain a fully conjugated system and thus

have very characteristic UV absorption profiles (Fig. 8) that make them easy to tentatively identify in an extract as a group. However, many variations with the same molecular formula make it difficult to make a positive identification on the basis of UV or MS data alone. Bicoumarins seems to be present in most *A. niger* strains, although they sometimes coelute with the aurasperones, which can obscure their detection. Orlandin, kotanin, and desmethyl-orlandin (based on LC-DAD, and LC-DAD-TOFMS) are consistently produced by *A. niger* and *A. tubingensis*. Another polyketide is funalenone [55], which is consistently (based on LC-DAD-TOFMS) produced by *A. niger* but also by *A. tubingensis* and *A. brasiliensis*.

The bicoumarins have been found to have some inhibitory effects on plant growth [56] and a moderate cytotoxic effect in human cell lines in vitro [57], but at present they are considered nontoxic [56, 59]. Yet, potential bioactivities of compounds of this type have not been investigated much and as a consequence no methods and data exist for their analysis.

Malformins

The malformins (Fig. 9) got their name owing to their ability to cause malformation in plants [60], and have also been reported to be antibacterial [61] and to inhibit interleukin-1 β binding to various human cells [62]. The malformins, which are cyclic pentapeptides with a sulfur bridge, have often been called mycotoxins since they have been shown to be toxic after peritoneal injection. However, since they were not toxic by oral administration [63] they cannot currently be considered mycotoxins *sensu stricto*.

In the *A. niger* group, only *A. brasiliensis*, *A. niger*, and *A. tubingensis* produce malformins [5, 64, 65], with *A. brasiliensis* probably being the best producer (confirmed by both LC-DAD-TOFMS and LC-MS/MS). Since they are produced by several abundant food contaminants, a

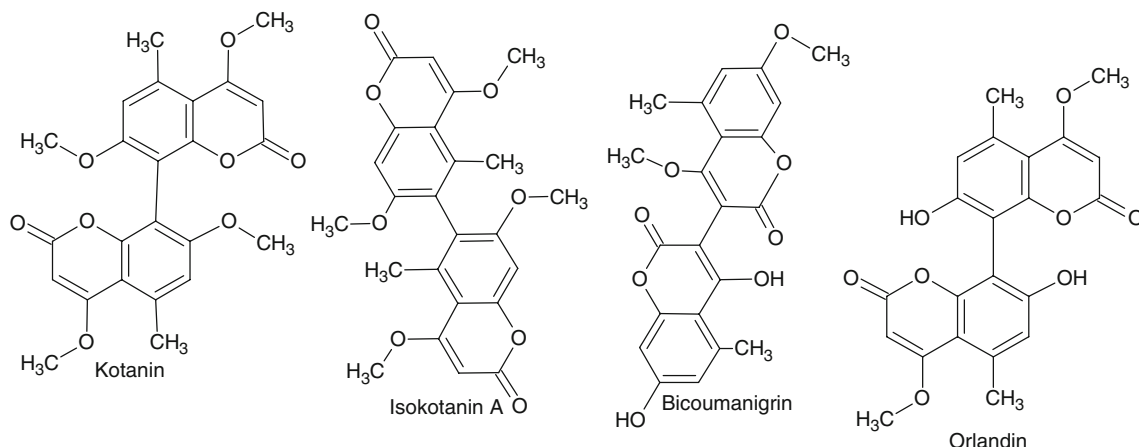


Fig. 7 Structure of selected bicoumarins

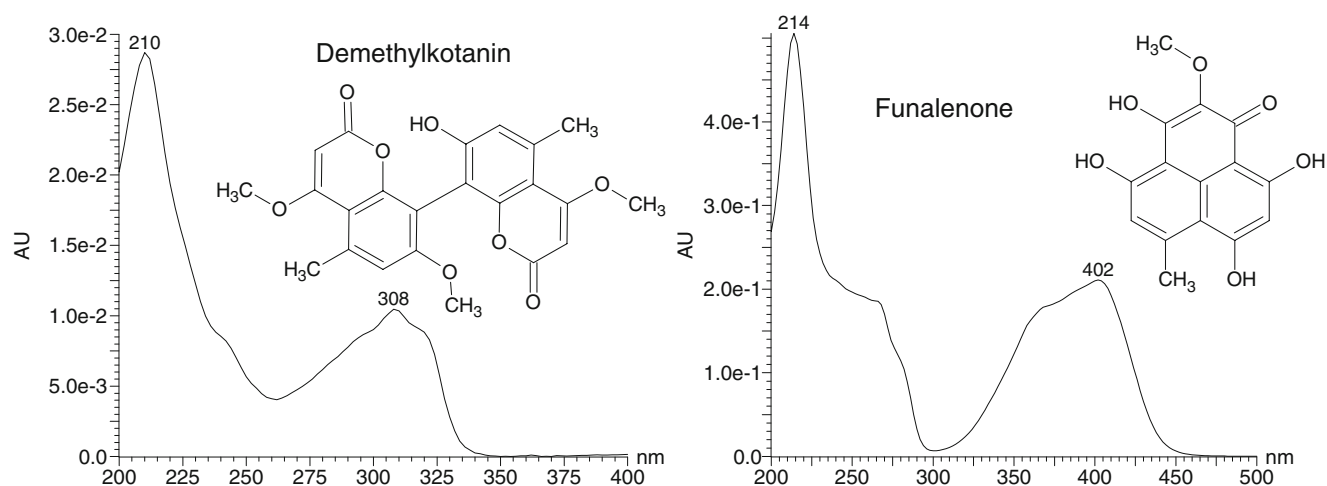


Fig. 8 Structure and UV spectrum of selected bicoumarins

toxicological evaluation, including bioavailability testing, is highly needed.

Even though malformins can be detected using LC-UV detection [19], in our experience they cannot be detected in *Aspergillus* extracts using this method owing to many interfering peaks. That is why LC-MS is the obvious choice for their detection [19, 66]. The sulfur bridge should also enable high-sensitivity electrochemical detection.

Chromatographic conditions used for analysis of the malformins are all based on acidic reversed chromatography [19, 66, 67], as normal phase and paper LC were not efficient [67]. Kim et al. [67] used a purification scheme based on acidic isocratic water–methanol (35:65 v/v) LC on an octadecyl silica column.

In ESI⁺, the malformins have a high tendency to form [M+NH₄]⁺, [M+Na]⁺, and [M+K]⁺ adducts [19], often with

the abundance of [M+H]⁺ 20% lower than that of the [M+NH₄]⁺ ion. If high in-source fragmentation settings are used, sodium and potassium adducts will dominate. In ESI⁺, *m/z* 417 is often observed from in-source fragmentation, and it is also a specific fragment for MS/MS analysis (MRM) along with *m/z* 372 and 304. In ESI[−], the malformins do not ionize as well as in ESI⁺ (tested on both MS instruments used) and are mainly detected as the [M+HCOO][−] and [M+Cl][−] adducts, with the abundance of [M-H][−] 20% lower than that of the other two adducts.

Asperazines

Asperazine (Fig. 10) is a complex diketopiperazine dimer, first isolated from a marine-derived *A. tubingensis* (reported as *A. niger*) by Varuglu et al. [64]. An analogue was later described by Ovenden et al. [68], also from an *A. tubingensis* strain. Asperazine was previously found to be cytotoxic in vitro against leukemia [64], but a later investigation reported it to be noncytotoxic [69].

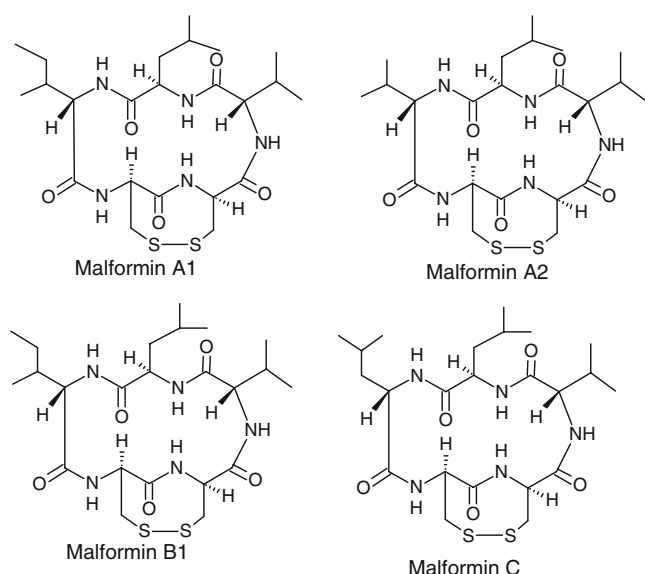


Fig. 9 Structures of selected malformins

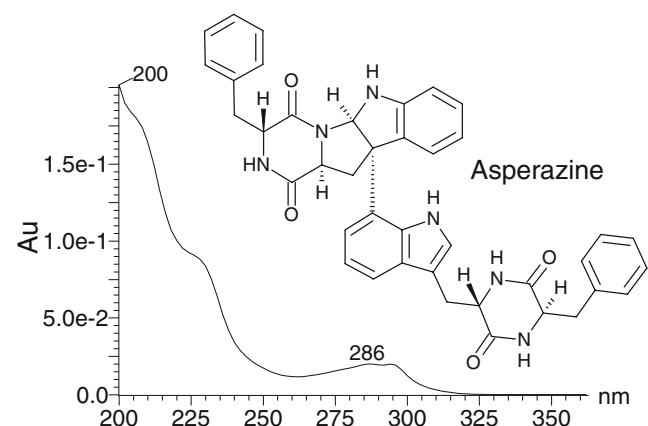


Fig. 10 UV spectrum and structure of asperazine

Reanalysis of extract data (approximately 66% by LC-TOFMS and all by LC-FLD) from 140 *A. niger*, 177 *A. tubingensis*, one *A. vadensis* [5], and 47 *A. acidus* (*A. foetidus*) [46, 70] strains for the production of asperazine showed that none of the *A. niger* strains produced asperazine, whereas a consistent production was observed in *A. acidus* and *A. tubingensis*. Owing to this limited distribution within the group, asperazine seems to be a valuable chemical marker that can be used to distinguish less toxic species such

as *A. acidus* and *A. tubingensis* from otherwise similar but more toxic species such as *A. niger*.

As most alkaloids, asperazines ionize very well in ESI⁺, with [M+H]⁺ as the only ion detected. Furthermore, asperazines show strong fluorescence at 230–450 nm (acidic conditions), and can usually also be detected from their distinct UV spectra using LC-DAD (UV/vis) (Fig. 10). Only acidic reversed-phase separations have been reported for their separation [5, 64, 70].

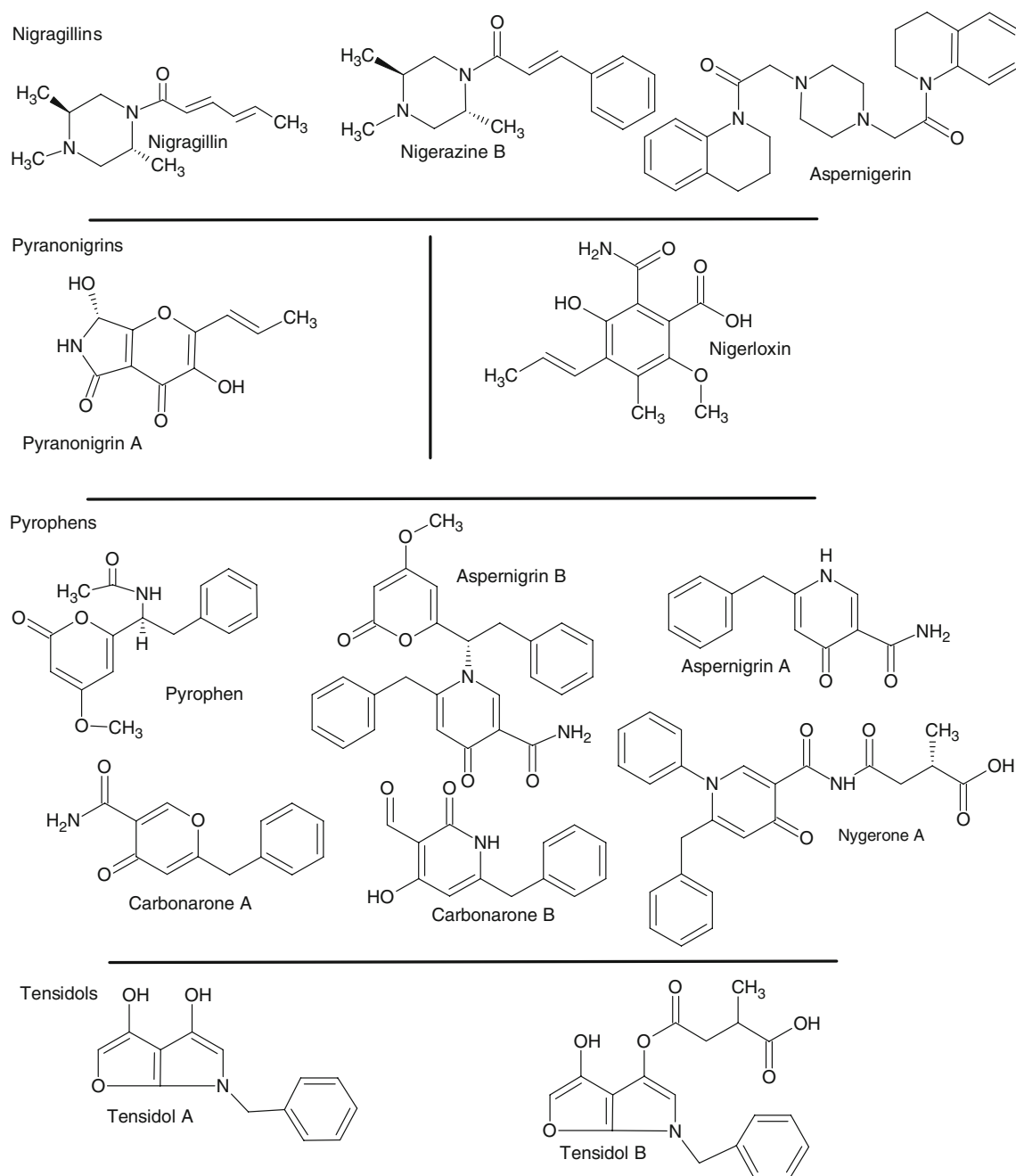


Fig. 11 Structures of the five classes of alkaloids

Other alkaloids

Species in the *A. niger* group produce a number of other nitrogen-containing compounds besides the fumonisins, ochratoxins, malformins, and asperazines. These alkaloids can be grouped into nigragillins, pyranonigrins, nigerloxin, pyrophens, and tensidols (Fig. 11).

Very few data are available for these compounds, with respect to both biological activities and which species they are found in. To our knowledge no studies exist on their presence in food and feeds, which is no surprise since they are, as is also the case for the NGPs, bicoumarins, and malformins, not commercially available.

Owing to their inherent properties as alkaloids, all of the compounds in this class have been found to ionize very well in ESI^+ and form strong $[\text{M}+\text{H}]^+$ ions, and except for the tensidols, they have very poor or no signal in ESI^- . In addition, many of them have characteristic chromophores suitable for LC-DAD.

The nigragillin-like group of alkaloids in the black aspergilli comprise nigragillin, nigerazines A and B [14, 71], aspernigrin B [57, 72], and aspernigerin [73]. Previous reports on nigragillin production within the *A. niger* group are restricted to *A. niger* (reported as *A. phoenicis*) [74]. Nigragillin was found on *A. niger* infected building materials, where the overall profile appeared similar to that of most rich-solid agar-based substrates [53].

We have found that nigragillin is easy to detect using LC-DAD-TOFMS, forming $[\text{M}+\text{H}+\text{CH}_3\text{CN}]^+$ besides $[\text{M}+\text{H}]^+$. Notably, to obtain retention in reversed-phase LC, the start gradient should not exceed more than 10% CH_3CN .

Results based on LC-DAD and LC-TOFMS show that all the species (but not all strains) in the “*A. niger* clade” produce pyranonigrins, mainly as pyranonigrin A [5]. The structure of pyranonigrin was first reported by Hiort et al. [57], who later revised it [58] and recently Schlingmann et al. [75] revised it again.

Pyranonigrin ionizes strongly in ESI^+ as $[\text{M}+\text{H}]^+$, with some $[\text{M}+\text{Na}]^+$ and $[\text{M}+\text{Na}+\text{CH}_3\text{CN}]^+$ adducts also being

formed. Owing to the relatively high polarity of these compounds, it is necessary to start a chromatographic gradient in the reverse phase at maximum 10% CH_3CN to obtain proper baseline separation from other early eluted compounds.

According to our data, pyrophens and tensidols are restricted to production in *A. tubingensis* and *A. niger*. On the basis of UV data, both produce several pyrophen analogues, all with the same ion pattern of $[\text{M}+\text{H}]^+$, some $[\text{M}+\text{Na}]^+$, and $[\text{M}+\text{NH}_4]^+$ adducts. The tensidols have consistently been detected in *A. niger* and *A. tubingensis* [15], where tensidol B is produced in much higher quantities than tensidol A. They ionize strongly in ESI^+ as $[\text{M}+\text{H}]^+$, with some $[\text{M}+\text{NH}_4]^+$ and $[\text{M}+\text{Na}+\text{CH}_3\text{CN}]^+$ ions. Tensidol A loses the 2-methylbutanedioic acid moiety to form fragment m/z 230. In ESI^- $[\text{M}-\text{H}]^-$ was observed along with some $[\text{M}-\text{H}+\text{H}_2\text{O}]^-$ which must be formed via hydrolysis of one of the double bonds.

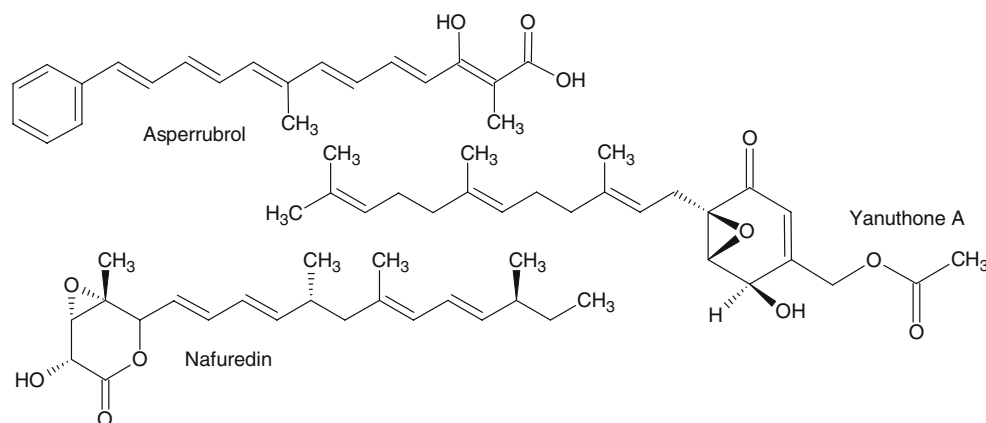
A novel analogue of pyrophen, nygerone A, was recently described by Henrikson [76] as a cryptic gene product which was induced by addition of histone inhibitors. Potentially, more new metabolites from *A. niger* or other black aspergilli could be unveiled by accessing cryptic gene clusters.

Remaining compounds

Very few terpenes have been isolated from the *A. niger* group, and they all contain a polyunsaturated chain as seen in Fig. 12. No data exist on the natural occurrence of terpenoids and investigation of biological activity has mainly been restricted to antimicrobial activities [77].

Reports of gibberellic acid [78] seem to be unlikely as the identity was only confirmed by a simple unspecific spectrometric assay. Another example is nigerloxin [79], where we have not found the compound in any extracts from the whole group when searching data files of approximately 150 extracts analyzed by LC-TOFMS for the $[\text{M}+\text{H}]^+$, $[\text{M}+\text{Na}]^+$, and $[\text{M}+\text{H}-\text{NH}_3]^+$ ions. The latter ion was included since screening of our reference standard

Fig. 12 Structure of selected terpenes



library has shown that NH_3 loss (using in-source fragmentation) is only observed from primary amides.

Biotransformation

This review is not intended to cover this area; however, it needs to be mentioned as some papers do not clearly state that the compounds detected from, for example, *A. niger* are actually biotransformed (small chemical changes to a complex molecule from other species) rather than fully biosynthesized by the fungus. Some of the well-known compounds derived from biotransformation are mentioned at the end of Table 2 and include compounds from animal feed or transformation of the feed material (genisteins, the flavonoids, isoflavone, pisolithin B, orobole), whereas iso-T-2 toxin is a transformation of the well-known *Fusarium* toxin T-2 toxin fed to the culture.

Analytical methods for mycotoxin screening

Summarizing the analytical observations, it appears that validated analytical methods only exist for OTA and the fumonisins. For the remaining compounds produced by the black aspergilli, the existing papers are mainly descriptions of the isolation procedure and structural elucidation, and no reference standards are commercially available. However, it should be possible to include them in an analytical scheme on a nonquantitative scale by using *A. niger* extracts for tuning MRM transitions when using triple quadrupole instruments, as, for example, is done for *F. avenaceum* metabolites in infected apples [80].

Profiles of culture extracts can also be compared with those of crude extracts of infected food and feed samples obtained using newer LC-TOFMS and Orbitrap instruments [41, 81]. These are getting increasingly more and more sensitive, and are a more interesting alternative to MRM analysis as they allow reprocessing of data files for more compounds [82]. With this approach, it is our recommendation to simultaneously monitor compounds within a series, for example, both FB_2 and FB_4 , several of the malformins (A–C), or several NGPs, as compounds within a class are usually always coproduced. This will help pinpoint suspicious samples, where, for example, only one malformin is detected.

Given that these compounds will be produced in infected crops, monitoring of NGPs, tensidols, fumonisins, malformins, bicoumarins (kotanins), and asperazines should enable the detection of growth of black aspergilli to species level.

Even though LC combined with DAD and accurate mass determination is a strong tool for identifying mycotoxins and other fungal metabolites, a correct elementary composition, characteristic UV spectra, and the same elution

profile [47, 83] are not sufficient for unambiguous identification of positional isomers of, for example, the NGPs or malformins. Detection in the original strain will provide a very strong tentative identification; however, for absolute identification NMR data are required. Improvements in the area of LC-NMR have made it an option even for the analytical chemist, as it is now possible to obtain data in the nanomole range [84]. Combining accurate mass determination with a few NMR recognizable features will make it possible to quickly identify positional isomers of known compounds [84].

In addition, it should be possible to quantify compounds in a fraction by NMR, since signals are proportional the number of moles in the tube. Subsequently one can calibrate against other (standardized) tubes containing accurate amounts of other compounds [85].

Conclusion

In conclusion, species within *Aspergillus* section *Nigri* are excellent producers of a large number of diverse secondary metabolites. Several new metabolites and maybe even new biosynthetic pathways are expected to be discovered in the near future now that the full genome of *A. niger* has been sequenced and soon also the full genomes of other black aspergilli will be sequenced.

Currently LC-DAD with accurate mass determination provides the easiest and most efficient strategy for tentative mapping of secondary metabolites in *A. niger* and its close relatives, especially if compared with already published chromatographic profiles, MS/MS data, and UV whole spectra. If absolute identification and positional isomer identification is needed, LC-NMR will be necessary.

For determination in food and feed, direct analysis of diluted crude extracts using LC-MS/MS analysis or LC-high-resolution mass-spectrometric detection is suggested. If sample pretreatment is needed, care should be taken when using anion exchange since *A. niger* can produce extremely high amounts of organic acids which can outsalt acidic target metabolites.

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